Sequence

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LENGTH: 22 amino acids
ryPR: amino acids
ryRANDEDNESS: single
rOPOLOGY: linear
MOLECULE TYPE: No. 6004925e
US-08-940-095-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
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                        STATE:
Sequence 30, Appl
Sequence 100, App
Sequence 4, Appli
Sequence 30, Appl
Sequence 100, App
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100, App
4, Appli
30, Appl
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4, Appli
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1, Appli
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Sequence 100, App
Sequence 4, Appli
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4, Appli
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                                                                                 August 16, 2007, 22:48:51; Search time 73 Seconds (without alignments) 43.032 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Description
                                                                                                                                                                                                                                                                                                                                                                 Issued Patents AA:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/5_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/6_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/7_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/H_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/H_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/RE_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/RE_COMB.pep:*
:: /EMC_Celerra_SIDS2/ptodata/1/iaa/RE_COMB.pep:*
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            GenCore version 6.2.1
Copyright (c) 1993 - 2007 Biocceleration Ltd.
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US-08-940-093-30

US-08-940-093-30

US-08-940-096-30

US-08-940-096-30

US-09-465-719-4

US-09-465-719-100

US-09-465-719-30

US-09-465-719-30

US-09-465-719-30

US-09-465-719-100

US-09-465-719-30

US-09-465-719-30

US-09-465-719-30

US-09-465-310

US-09-453-838-4

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US-09-453-838-30
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US-09-453-841-100
US-09-453-833-4
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1 PVLDLFRELLNELLEALKQKLK 22
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Maximum Match 100%
Listing first 45 summaries
                                                              OM protein - protein search, using sw model
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Maximum DB seq length: 200000000
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Match
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Perfect score:
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                                                                                                                                                                  Sequence:
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No.
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4, Appli
30, Appl
100, App
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30, Appl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Sekul, Renate
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
APPLICANT: Dufourcq, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-I ACONISTS
NUMBER OF SEQUENCES: 258
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                       Sequence
Sequence
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US-09-453-833-30
US-09-453-823-100
US-09-453-826-4
US-09-453-826-100
US-09-453-826-100
US-09-453-840-4
US-09-453-840-100
US-09-453-840-100
US-09-865-989-4
US-09-865-989-100
US-09-865-989-100
US-09-865-989-100
US-09-453-834-100
US-09-453-834-100
US-10-283-599-30
US-10-283-599-30
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US-09-465-718-30
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: STATEM 'Version 2.0
SOFTWARE: FESTERN' Oversion 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
CLASSIPICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/08940095
Patent No. 6004925
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Coruzzi, Laura A
REGISTRATION UNDBER: 30,742
REFERENCE/DOCKET NUMBER: 0091
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELERX: 66141 PENNIE
TELEX: 66141 PENNIE
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NAME/KEY: Other
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    JS-08-940-095-100
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                                                                                                                                             Sequence 30, Application US/08940095

Sequence 30, Application US/08940095

Patent No. 6004925

GENERAL INFORMATION:

APPLICANT: Desseux, Jean-Louis

APPLICANT: Sekul, Renate

APPLICANT: Cornut, Isabelle

APPLICANT: Cornut, Isabelle

APPLICANT: Metz, Gunther

APPLICANT: Dufourcq, Jean

TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS

TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS

NUMBER OF SEQUENCES: 258

CORRESPONDENCE ADDRESS:

ADDRESSEE: Pennie & Edmonds LLP

STREET: 1155 Avenue of the Americas

CITY: New York

STATE: NY
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Query Match

100.0%; Score 105; DB 2; Length 22;
Best Local Similarity 100.0%; Pred. No. 4.9e-07;
Matches 22; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1...22
CTHER INFORMATION: N-terminal dansylated peptide US-08-940-095-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
CLASSIFTCATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: BY
FILING DATE:
APPLICATION NUMBER: 30,742
RESISTRATION NUMBER: 30,742
RESISTRATION NUMBER: 30,742
RESISTRATION NUMBER: 30,742
RESISTRATION NUMBER: 30,742
RESERNICE/DOCKET NUMBER: 009196-0004-999
TELEPHONE: 650-493-4935
                                                                                           1 PVLDLFRELLNELLEALKOKLK 22
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: No. 6004925e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
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ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Other
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PVLDLFRELLNELLEALKOKLK 22

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APPLICANT: Dasseux, Jean-Louis
APPLICANT: Betul. Renate
APPLICANT: Setul. Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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CTHER INFORMATION: All amino acids are in the D-configuration US-08-940-095-100
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100.0%; Pred. No. 4.9e-07;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  009196-0004-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IEM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FESTENCY O'ERSION 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
CLASSIPICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/08940093; Patent No. 6037323; GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis APPLICANT: Sekul, Renate APPLICANT: Buttner, Klaus APPLICANT: Cornut, Isabelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PVLDLFRELLNELLEALKOKLK 22
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sekul, Renate
Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: No. 6004925e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.
Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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NAME/KEY: Other
LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 105; DB 2; 100.0%; Pred. No. 4.9e-07;
                          PILING DATE: 29-SEP-1997
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER:
PILING DATE:
PILING DATE:
ATORNEY/AGENT INFORMATION:
NAME: COLUZZI, LBULTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
RELERRANICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELERRAN: 650-493-5556
TELERY: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,093
FILING DATE: 29-SRE-1997
CLASSIFICATION WHEE:
APPLICATION WHEE:
RICH APPLICATION DATA:
APPLICATION WHEE:
ALPHICATION WHEE:
                        US/08/940,093
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                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: No. 6037323e
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Matches 22; Conservative
CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
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COUNTRY:
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GENERAL INFORMATION:

APPLICANT: Baseux, Jean-Louis
APPLICANT: Sekul, Renate

APPLICANT: Sekul, Renate

APPLICANT: Genther

TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS

TITLE OF INVENTION: APOLIPOPROTEIN B-I AGONISTS

TITLE OF SEQUENCES: 258

CORRESPONDENCE ADDRESS:

ADDRESSE: Pennie & Edmonds LLP

STREET: 1155 Avenue of the Americas

CITY: New York

STATE: NY
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                              COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: DISKETE
COMPUTER: DISKETE
COPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,093
FILING DATE: 29-SEP-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 05-SEP-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 30,742
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECOMMUNICATION INFORMATION:
TELEPRAX: 66141 PENNIE
TELEPRAX: 66141 PENNIE
TELEPRAX: 66141 PENNIE
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TVENEY. A mino acids
TVENEY. A mino acids
                                          NUMBER OF SEQUENCES: 258
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
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Patent No. 6037323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. 6037323e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-940-093-30
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Length 22; Indels

Gaps

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Length 22; Indels

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APPLICANT: Setul, Render APPLICANT: Setul, Render APPLICANT: Buttner, Klaus APPLICANT: Buttner, Klaus APPLICANT: Buttner, Klaus APPLICANT: Gornut, Isabelle TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS CORRESPONDENCE ADDRESS: ADDRESSE: Pennie & Edmonds LLP STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 105; DB 2;
100.0%; Pred. No. 4.9e-07;
tive 0; Mismatches 0;
                                                                          100.0%; Score 105; DB 2; 100.0%; Pred. No. 4.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        009196-0005-999
                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM COMPUTER:
COMPUTER: IBM COMPUTER:
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,096
                                                                                                                                                                                                                                                 1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                           1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30, Application US/08940096
Patent No. 6046166
GENERAL INFORMATION:
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ATTORNEY AGENT INFORMATION:
NAME: COTUZZI, LGUTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00915
TELECOMMUNICATION INFORMATION:
TELECHONE: 650-493-5556
TELECAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 22 anino acids
LENGTH: 22 anino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: No. 6046166e
                                            Query Match
Best Local Similarity 100.v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 29-SEP-1
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
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          US-08-940-096-4
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APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
MANDERS
                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Other
1 LOCATION: 1...22
1 COTATION: All amino acids are in the D-configuration US-08-940-093-100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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; Pred. No. 4.9e-07;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRICE APPLICATION BATA:
APPLICATION NATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY AGENT INFORMATION:
NAME: COLUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0005-999
TELECOMMUNICATION INFORMATION:
TELEFRAX: 650-493-4935
TELEFAX: 650-493-5556
TELEFAX: 6614 PENNER
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,096
FILING DATE: 29-SEP-1997
CLASSIFICATION: 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/08940096
Patent No. 6046166
GENERAL INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100: SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: linear
TOPOLOGY: linear
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%;
Best Local Similarity 100.0%;
Matches 22; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: No. 6046166e
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ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JS-08-940-096-4
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Gaps

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Indels

Length 22;

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APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
CORRESPONDENCES: 258
CORRESPONDENCE ADDRESS:
TITLE OF INVENTION: AND THEIR USB TO TREAT DYSLIPIDEMIC DISORDERS NUMBER OF SEQUENCES: 258
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 105; DB 2; Length 22; 100.0%; Pred. No. 4.9e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/465,719
                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,093
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PVLDLFRELLNELLEALKQKLK 22
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; Sequence 30, Application US/09465719
; Patent No. 6265377
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dasseux, Jean-Louis
Sekul, Renate
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TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: No. 6265377e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM COMPACTIBLE OPERATING SYSTEM: DOS SOFTWARE: FASTSEO Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELERAX: 650-499-5556
TELEX: 66141 PENNIE
INFORMATION POR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036-2811
COMPUTER READABLE FORM
MEDIUM TYPE: Disket
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
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                                        Sequence 100, Application US/08940096
Fatent No. 6046166
GENERAL INFORMATION:
APPLICANT: Beature, Klaus
APPLICANT: Buttner, Klaus
APPLICANT: Buttner, Klaus
APPLICANT: Metz, Gunther
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
ITILE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1...22
CTHER INFORMATION: All amino acids are in the D-configuration US-08-940-096-100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Buttner, Klaus
APPLICANT: Merz, Gunther
ITLE OP INVENTION: APOLIPOPROTEIN A-1 AGONISTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDING TYPE DISKETCE
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastERO Version 2.0
CURENY APPLICATION DATA:
APPLICATION NUMBER: US/08/940,096
FILING DATE: 29-SEP-1997
CLASSIFICATION NUMBER: US/08/940,096
FILING DATE: APPLICATION DATA:
APPLICATION NUMBER: 30
FILING DATE: APPLICATION NUMBER: 30,742
FEGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0005-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEPHONE: 650-493-5556
TELERAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 22-gming acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: No. 6046166e
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Other
                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                 -08-940-096-100
                                                                                                                                                                                                                                                                                                                                                                                                        STATE: NO COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g
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Gaps

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TELEFAK: 650-493-5556
TELEK: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 66141 PENNIE
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                650-493-5556
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TELEPHONE:
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STATE:
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Patent No. 6265377

GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Butcher, Klaus
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: APPLIPEN USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          i LOCATION: 1...22
; OTHER INFORMATION: N-terminal dansylated peptide
US-09-465-719-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 105; DB 2; Best Local Similarity 100.0%; Pred. No. 4.9e-07; Matches 22; Conservative 0; Mismatches 0;
                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,093
FILING DATE: 29-SEP-1997
ATTORNEY AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELEPHONINICATION INFORMATION:
TELEPHONE: 650-493-5556
TELEPAX: 650-493-5556
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/940,093
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: COTULATI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/465,719
APPLICATION NUMBER: US/09/465,719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: No. 6265377e
                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie &
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COMPUTER READABLE FORM:
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                   FILING DATE:
CLASSIFICATION:
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| INTROMMENTOR FOR ENGINE
| SEGURATES | SEGURATES |
| STANDEDWESS | SEGURATES |
| TOPOLOGY: 1 | Interest |
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US-09-453-605-30
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                                                                                                                                                                                                                                    Sequence 4, Application US/09453605
Sequence 4, Application US/09453605
Patent No. 6329341
GENERAL INPORMATION:
Sekul, Renate
Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
Dufourcq, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: No. 6329341e
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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Patent No. 6329341
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 PVLDLPRELLNELLEALKOKLK 22
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1 PVLDLFRELLNELLEALKOKLK 22
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TELEPHONE: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Buttner, Klaus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
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                                                                                                                                                                                                                  US-09-453-605-4
                                                                                                                                                                      RESULT 14
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Cornut, Isabelle
Metz, Gunther
Dufourcq, Jean
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPDEMIC DISORDERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Other LOCATION: 1...22 OTHER INFORMATION: N-terminal dansylated peptide SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 009196-0004-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 105; DB 2; Best Local Similarity 100.0%; Pred. No. 4.9e-07; Matches 22; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                    CONNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PEALSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/453,605
FILING DATE: 26-NO. 6329341-1999
CLASSIFICATION: <a href="https://doi.org/10.1007/">chassification</a>
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: 08/940,095
                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Search completed: August 16, 2007, 22:50:33
Job time : 74 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: No. 6329341e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 650-493-5556
TELEX: 66141 PENNIR
INFORMATION FOR SEQ 1D NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FOPOLOGY: linear
                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                              CITY: New York
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Appli Appli

Sequence Sequence

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Sequence

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Sequence Sequence

Sequence Sequence Sequence

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Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
AND THEIR USE TO TREAT DYSLIPEDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 30,742
TELECOMMUNICATION INFORMATION:
TELERPHONE: 650-493-4935
               US-10-099-574A-4

US-10-099-574A-30

US-10-099-574A-100

US-10-715-895-30

US-11-715-895-30

US-11-482-292-100

US-11-482-292-30

US-11-482-292-100

US-11-482-999-7

US-11-683-784-1

US-09-865-989-7

US-10-099-574A-7

US-10-099-574A-7

US-10-099-836B-7

US-11-683-784-1

US-10-099-836B-7

US-10-099-836B-7

US-10-099-836B-7

US-10-099-836B-7

US-10-099-836B-7
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ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/865,989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: NO. US20030008827Ale SEQUENCE DESCRIPTION: SEQ ID NO: 4: US-09-865-989-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 25-May-2001
CLASSIFICATION: cUnknown>
PRIOR APPLICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/0965989
Publication No. US20030008827A1
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 22 amino acids
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 258
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2: /EMC_Celerra_SIDS2/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
3: /EMC_Celerra_SIDS2/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
4: /EMC_Celerra_SIDS2/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
5: /EMC_Celerra_SIDS2/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
6: /EMC_Celerra_SIDS2/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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                                                                                                                                                                  August 16, 2007, 22:49:31 ; Search time 300 Seconds (without alignments) 59.976 Million cell updates/sec
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                      GenCore version 6.2.1
Copyright (c) 1993 - 2007 Biocceleration Ltd.
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US-10-099-574A-4
US-10-099-574A-30
US-10-099-574A-100
US-10-25-940-1
US-10-099-836B-4
US-10-099-836B-30
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US-10-283-599-4
US-10-283-599-30
US-110-283-599-100
US-110-802-080-4
US-110-802-080-30
US-110-802-080-100
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Post-processing:

Database

Result No.

Title: Perfect score:

Sequence:

Scoring table:

Searched:

us-10-715-895a-4.rapbm

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US-09-865-989-4
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Cornut, Isabelle
Metz, Gunther
TITLE OF INVENTION AND THEIR USE TO TREAT DYSLIPDIBMIC DISORDERS
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                Score 105; DB 3; Length 22; Pred. No. 1.8e-07; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMER: 30,742
REFERENCE/DOCKET NUMBER: 309196-0006-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFRAX: 650-493-5556
TELEFRAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/865,989
FILING DATE: 25-May-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: No. US20030008827A1e
                                                                                                                                                                                         US-09-865-989-30; Sequence 30, Application US/09865989; Publication No. US2003008827A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                           1 PVLDLFRELLNELLEALKOKLK 22
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LENGTH: 22 amino acids
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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STRANDEDNESS: single
                100.0%;
ilarity 100.0%;
Conservative 0
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Matches 22; Conservative
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Ouery Match
Best Local Similarity
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                                                                                                                    Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-1 AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
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APPLICATION NUMBER: US/09/865,989
FILING DATE: 25-May-2001
CLASSIFICATION NUMBER: US/09/865,989
PRIOR APPLICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORNEY/AGENT INFORMATION:
NAME: COTULET, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECHONEY 650-493-4935
TELEPHONE: 650-493-5556
TELERY: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
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Cornut, Isabelle
Metz, Gunther
Hetz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
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100.0%; Pred. No. 1.8e-07;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. US20030008827Ale
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 2.0
Sequence 100, Application US/09865989
Publication No. US20030008827A1
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/0965989
Publication No. USZO040029807A9
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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Matches 22; Conservative
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CURRENT APPLICATION DATA:
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Best Local Similarity
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Cornut, Isabelle
Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
  AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                               CONPUTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER IBM COMPALIDIE
OPERATING SYSTEM: DOS
SOFTWARE: FASISEQ Version 2.0
CURRENT APPLICATION NUMBER: US/09/865,989
FILING DATE: 23-May-2001
CLASSIFICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORICATION NUMBER: 30,742
REFIRENCE COLUZZI, LAULA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELEPRONE 650-493-4935
TELEPRONE: 650-493-556
TELERY: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TVOME: amino acids
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                            CORRESPONDENCE ADDRESS:
ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: No. US20040029807A9e
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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US-09-865-989-30
Sequence 30, Application US/09865989
Publication No. US20040029807A9
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Sexul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLDLFRELLNELLEALKOKLK 22
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ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                       NUMBER OF SEQUENCES: 258
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Sequence 100, Application US/09865989
Publication No. US20040029807A9
GENERAL INFORMATION:
APPLICANT: Basseux, Jean-Louis
Sekul, Renate
Buttner, Klaus
Cornut, Isabelle
Metz, Gunther:
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
APPLICATION NUMBER: US/09/865,989
FILING DATE: 25-May-2001
CLASIPICATION: cUbknown>
PRIOR APPLICATION DATE:
APPLICATION OWHER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORNEY, AGENT INFORMATION:
NAME: CORUZZI, LAULE A
REGISTRATION NUMBER: 30,742
REPERENCE, DOCKET NUMBER: 009196-0006-999
TELEPHONICATION INFORMATION:
TELEPHONIE: 650-493-4955
TELEFRX: 650-493-4955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 105; DB 3; 100.0%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEO Version 2.0
CURRENT APPLICATION DATA:
CLASSIPICATION: CURREN: US/09/865,989
FILING DATE: 25-May-2001
CLASSIPICATION: CURROWN:
PRIOR APPLICATION: CURROWN:
APPLICATION: AURION:
APPLICATION NUMBER: 09/465,719
FILING DATE: 17-DEC-1999
ATTORNEY/AGENT INFORMATION:
MAMB: COTUZZI, LAURA A
REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSER: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: No. US20040029807A9e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                            TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30.
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Other LOCATION: 1...2
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TOPOLOGY: linear MOLECULE TYPE: No. US20030060604Ale
                                 ;
US-10-099-574A-4
                                                                                          Query Match
                                                                                                                                                                                                                                                                                          RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                               NAME/KEY: Other LOCATION: 1...22 OTHER INFORMATION: All amino acids are in the D-configuration SEQUENCE DESCRIPTION: SEQ ID NO: 100:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/10099574A

Publication No. US20030060604A1

GENERAL INFORMATION:
APPLICANT: Basseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Gornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-1 AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 254
CORRESPENDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 105; DB 3; Length 22; Best Local Similarity 100.0%; Pred. No. 1.8e-07; Matches 22; Conservative 0; Mismatches 0; Indels
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-5556
TELEPAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               009196-0005-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/099,574A
FILING DATE: 29-SEP-1997
CLASSIFICATION:
                                                                                                                                                                           TYPE: amino acid
STRANDENESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. US20040029807A9e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PVLDLFRELLNELLEALKQKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: COLUZZI, LAUER A
REGISTRATION NUMBER: 30,742
REPRENCE/DOCKET NUMBER: 0091:
TELECOMMUNICATION INFORMATION:
TELEPAN: 650-493-4935
TELERA: 66141 PENNIE
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                   TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
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STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                         US-09-865-989-100
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US-10-099-574A-4
                                                                                                                                                                                                                                                                    FEATURE
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APPLICANT: Bactul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & PA---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                       Gaps
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100.0%; Score 105; DB 4; Length 22; 100.0%; Pred. No. 1.8e-07;
                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 105; DB 4; 100.0%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               009196-0005-999
                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PASISEO VERSION 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/099,574A
FILING DATE: 29-SEP-1997
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
FILING DATE:
ATORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: No. US20030060606Ale
                                                                                                                                                                                       US-10-099-574A-30
; Sequence 30, Application US/10099574A
; Publication No. US20030060604A1
; Publication No. US20030060604A1
; APPLICANT: Dasseux, Jean-Louis
                                                                                                1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PVLDLFRELLNELLEALKOKLK 22
                                                                        1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFRENCE/DOCKET NUMBER: 009:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 aming acids
                                     Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.
Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: NY
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                                                                                                                         APPLICANT: Dasseux, Jean-Louis
APPLICANT: Batenate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Cornut, Isabelle
APPLICANT: Many APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF ADDRESS:
CORRESPONDENCES: 254
CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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1.00ATION: 1...22

COLATION: All amino acids are in the D-configuration US-10-099-574A-100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/10252940
; Publication No. US20030099714A1
GENERAL INPORMATION:
; TITLE OF INVENTION: PEPTIDE/LIPID COMPLEX FORMATION
BY CO-LYOPHILIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     009196-0005-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/099,574A
FILING DATE: 29-SEP-1997
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: No. US20030060604Ale
                               Sequence 100, Application US/10099574A Publication No. US20030060604A1 GENERAL INFORMATION: APPLICANT: Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PVLDLFRELLNELLEALKOKLK 22
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APPLICATION NUMBER:
PILING DATE:
ATORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
REPRENCE/POCKET NUMBER: 00919
TELECOMMUNICATION INFORMATION:
TELERAX: 650-493-4935
TELEFAX: 650-493-5556
TE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 22 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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JS-10-099-574A-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-252-940-1
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Butther, Klaus
Cornut, Isabelle
Metz, Gunther
Dufourcg, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-1 AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                              COMPUTER: ITES CARACTES
CONTURED: FactStore
SOFTWARE: FactStore
CURRENT APPLICATION DATA:
CLASSIFICATION NUMBER: US/10/252,940
FILING DATE: 23-Sep-2002
CLASSIFICATION NUMBER: US/08/942,597
FILING DATE: 02-OCT-1997
APPLICATION NUMBER: US/08/942,597
FILING DATE: 02-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 9196-0008-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELERPHONE: 650-493-556
TELERPHONE: 650-493-556
TELER: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 105; DB 4;
100.0%; Pred. No. 1.8e-07;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                         ADDRESSEE: Pennie & Edmonds, LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
CURRENT ADDATA: APPLICATION NUMBER: US/10/099,836B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10099836B
Publication No. US20030203842A1
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 22 amino acids
                                                                                                             COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 22; Conservative
                                                                     CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 11
US-10-099-836B-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-252-940-1
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Gaps

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Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
Dufourcg, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-1 AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                          NAME/KEY: Other
LOCATION: 1...22
OTHER INFORMATION: N-terminal dansylated peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: DISACLE
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOCTHWARE: FRESESQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/099,836B
FILING DATE: 28-Aug-2002
CLASSIFICATION: «Unknown»
FILING DATE: Andrown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: «Unknown»
FILING DATE: «Unknown»
ATORNEY/AGENT INFORMATION:
NAME: COTUZEL, LABUE A
REGISTRATION NUMBER: 30,742
RESERBENGE/BOCKET NUMBER: 009196-0004-999
TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 105; DB 4;
Best Local Similarity 100.0%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                            TOPOLOGY: linear
MOLECULE TYPE: No. US20030203842Ale
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: No. US20030203842Ale FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 13
US-10-099-836B-100
i Sequence 100, Application US/10099836B
i Publication No. US20030203842A1
i GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLDLFRELLNBLLEALKOKLK 22
                        SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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INFORMATION FOR SEQ ID NO: 100:
                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                           22; Conservative
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Publication No. US20030203842A1
GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
Butner, Klaus
Cornut, Isabelle
Metz, Gunther
Diffured, Jean
TITLE OF INVENTION: APDLIFOPROTEIN A-1 AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA

ZIP: 10036-2811

COMPUTER READABLE FORM:

WEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

COMPUTER: IBM Compatible

COMPUTER: STSTEM: DOS

SOFTWARE: FASTSO Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/099,836B

FILING DATE: 28-Aug-2002

CLASSIFICATION: CURKNOWN:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: CURKNOWN:

RILING DATE: CURKNOWN:

APPLICATION NUMBER: CURKNOWN:

RILING DATE: CURKNOWN:

APPLICATION NUMBER: CURKNOWN:

RILING COLUZZI, Lautra A

REGISTRATION NUMBER: 30,742

REFERENCE/DOCKET NUMBER: 009196-0004-999

TELLECOMMUTCATION:

REFERENCE/DOCKET NUMBER: 009196-0004-999
                                                                                       FILING DATE: «Unknown»
ATTORNEY/AGENT INPORMATION:
NAME: COTUZZI, Lauta A
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 009196-0004-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear

MOLECULE TYPE: No. US20030203842Ale

SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-10-099-836B-4
                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 PVLDLFRELLNELLEALKOKLK 22
                          CLASSIFICATION: <Unknown:
FILING DATE: 28-Aug-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                  TELEFAX: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                              FYPE: amino acid
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US-10-283-599-30
                         RESULT 15
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0
                                                                                                                                                            ö
LOCATION: 1...22 OTHER INFORMATION: All amino acids are in the D-configuration SEQUENCE DESCRIPTION: SEQ ID NO: 100:
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sakul, Fenate
APPLICANT: Sakul, Fenate
APPLICANT: Sakul, Fenate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Cornut, Isabelle
APPLICANT: Cornut, Isabelle
TITLE OF INVENTION: GENE THERAPY APPROACHES TO
TITLE OF INVENTION: USE TO TREAT DYSLIPIDEMIC DISORDERS.
NUMBER OF SEQUENCES: 274
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie F. F.
STREEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                            Gaps
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                                                                                                            Query Match 100.0%; Score 105; DB 4; Length 22; Best Local Similarity 100.0%; Pred. No. 1.8e-07; Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/POCKET NUMBER: 009196-0007-999
TELECOMMUICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/283,599
FILING DATE: 29-OCT-2002
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,136
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear _ MOLECULE TYPE: No. US20030208059Ale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                  1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                        1 PVLDLFRELLNELLEALKQKLK 22
                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/10283599
Publication No. US20030208059A1
GENERAL INFORWATION:
APPLICANT: Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 650-493-5556
TELEFX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                  US-10-099-836B-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                     RESULT 14
US-10-283-599-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-283-599-4
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PVLDLFRELLNELLEALKOKLK 22

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Sequence 30, Application US/10283599
; Publication No. US20030208059A1
; Publication No. US20030208059A1
; GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Cornut, Isabelle
APPLICANT: Grante
TITLE OF INVENTION: GENE THERAPY APPROACHES TO
TITLE OF INVENTION: GENE THERAPY APPROACHES TO
TITLE OF INVENTION: USE TO TREAT DYSLIPIDEMIC DISORDERS.
NUMBER OF SEQUENCES: 274
CORRESCONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: 1...2
; OTHER INFORMATION: N-terminal dansylated peptide
US-10-283-599-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 105; DB 4; 100.0%; Pred. No. 1.8e-07;
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                                                                                                                                                                                                                                                                                                               ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FRAESQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/283,599
FILING DATE: 29-OCT-2002
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: No. US20030208059Ale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,136
FILING DATE: 29-SBP-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLDLFRELLNELLEALKOKLK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 005
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity luv...
Best Local 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: single
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# November 2005

generate two sets of results each. The Published\_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being Published\_Applications Nucleic Acid and Published\_Applications Amino Acid database searches now available for processing searches. Newly published applications will appear in the Published\_Applications\_New databases; older published applications make up the Published Applications Main databases. Searches run against Nucleic Acid Published\_Applications produce two sets of results, with the extensions Searches run against Amino Acid Published\_Applications produce two sets of results, with the extensions .rnpbm (Published\_Applications\_NA\_Main) and .rnpbn (Published\_Applications\_NA\_New). .rapbm (Published\_Applications\_AA\_Main) and .rapbn (Published\_Applications\_AA\_New).

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rype: PRT GAGANISM: Homo sapien US-10-533-069-1050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-533-069-452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 900
셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24, Appl
116, App
23, Appl
1316, Ap
121, App
943, App
100, App
23, Appl
192, Appl
192, App
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                          August 16, 2007, 22:50:51; Search time 7 Seconds (without alignments) 3.121 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Description
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Sequence 3
Sequence 1
Sequence 2
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Sequence
Sequence
                   GenCore version 6.2.1
Copyright (c) 1993 - 2007 Biocceleration Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-533-069-1050
US-10-533-069-452
US-10-533-069-1216
US-10-533-069-1039
US-10-533-069-1039
US-10-533-069-1043
US-10-533-069-1043
US-10-533-069-1154
US-10-533-069-1156
US-10-533-069-1156
US-10-533-069-1156
US-10-533-069-1156
US-11-657-313-24
US-11-657-313-24
US-11-657-313-23
US-11-657-313-069-100
US-11-539-856-23
US-11-725-235-116
                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                            105
1 PVLDLFRELLNELLEALKOKLK 22
                                                                                               OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                             5387 segs, 993155 residues
                                                                                                                                                                                                                                                                                    BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                          US-10-715-895A-4
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2000
2000
11154
11154
1427
1427
1427
11350
2035
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Sequence 1050, Application US/10533069

Publication No. US20070186017A1

GENERAL INFORMATION:

APPLICANT: AGGARWAL, SUDEPTA

APPLICANT: GURNEY, AUSTIN L.

APPLICANT: SCHÖENFELD, JILL

APPLICANT: WILLIAMS, P. MICKEY

APPLICANT: WILLIAMS, P. MICKEY

APPLICANT: WILLIAMS, P. MICKEY

APPLICANT: WILLIAMS, P. MICKEY

APPLICANT: WU, THOWAS D.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE

TITLE OF INVENTION: RELATED DISEASES

FILE REFERENCE: P1962R1 US

CURRENT APPLICATION NUMBER: US/10/533,069

CURRENT APPLICATION NUMBER: PCT/US03/34381

PRIOR APPLICATION NUMBER: US 60/422,472

PRIOR FILING DATE: 2003-10-29

NUMBER OF SEQ ID NOS: 2442

SEQ ID NO 1050

TENGTH. ON 01050
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236, App
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860, App
1015, Ap
292, App
1600, Ap
725, App
855, App
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1172, Ap
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860, A
1015,
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42.1%; Pred. No. 13;
tive 7; Mismatches 4; Indels
                               US-10-533-069-735
US-10-533-069-860
US-10-533-069-1015
US-11-551-744-292
US-10-533-069-1600
US-10-533-069-1600
US-10-533-069-128
US-10-533-069-1172
US-10-533-069-1172
US-10-533-069-1976
US-10-533-069-1976
US-11-656-389-74
US-11-656-389-15
US-11-656-389-15
US-11-656-389-15
US-11-656-389-15
US-11-656-389-15
US-11-656-389-15
US-11-656-389-17
US-11-656-389-182
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Publication No. US20070185017A1
GENERAL INFORMATION:
APPLICANT: AGGARMAL, SUDEEPTA
APPLICANT: CLARK, HILARY
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: SCHOENPELD, JILL
APPLICANT: WILLIAMS, P. MICKEY
APPLICANT: WOOD, WILLIAM I.
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Best Local Similarity 42.1
Matches 8; Conservative
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Gaps

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APPLICANT: GURNEY, AUSTIN L.
APPLICANT: SCHOENFELD, JILL
APPLICANT: WILLIAMA;
PRELICANT: WILLIAMA;
APPLICANT: WILLIAMA;
APPLICANT: WU, THOMAS D.
TITLE OF INVENTION: COMPOSITIONS.AND METHODS FOR THE TREATMENT OF IMMUNE
TITLE OF INVENTION: RELATED DISEASES
FILE REFERENCE: P1982R1 US
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2003-10-28
FRIOR APPLICATION NUMBER: US 60/422,472
FRIOR PILING DATE: 2003-10-28
FRIOR PILING DATE: 2002-10-29
FRIOR PILING DATE: 2002-10-29
WHINER OF SEQ ID NOS: 2442
SEQ ID NO 1039
                                                             TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE TITLE OF INVENTION: RELATED DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 1154; 46;
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                                                                                                           CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2005-04-28
FRIOR APPLICATION NUMBER: PCT/US03/34381
FRIOR APPLICATION NUMBER: PCT/US03/34381
FRIOR FILING DATE: 2003-10-28
FRIOR APPLICATION NUMBER: US 60/422,472
FRIOR PILING DATE: 2002-10-29
FRIOR PILING DATE: 2002-10-29
FRIOR SEQ ID NOS: 2442
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Pred. No.
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Publication No. US20070185017A1
GENERAL INFORMATION:
APPLICANT: AGGARMAL, SUDEEPTA
APPLICANT: CLARK, HILARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1070, Application US/10533069
Publication No. US2070185017A1
Publication No. US2070185017A1
APPLICANT: AGGARWAL, SUDEEPTA
APPLICANT: CLARK, HILARY
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: SCHOENFELD, JILL
APPLICANT: SCHOENFELD, JILL
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109 EMLDDLMEKRKEKL 122
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Best Local Similarity 50.0
Matches 7; Conservative
   WOOD, WILLIAM I.
                                   WU, THOMAS D.
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; ORGANISM: Homo sapien
US-10-533-069-1039
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapien
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   APPLICANT:
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APPLICANT: CLARK, HILARY
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: WILLIAMS, P. MICKEY
APPLICANT: WOLLIAMS, P. MICKEY
APPLICANT: WOLLIAMS D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE
TITLE OF INVENTION: RELATED DISEASES
FILE REFERENCE: P1982R1 US
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2005-04-28
PRIOR APPLICATION NUMBER: PCT/US03/34381
PRIOR APPLICATION NUMBER: US 60/422,472
PRIOR PILING DATE: 2002-10-29
PRIOR FILING DATE: 2002-10-29
SEQ ID NO 1216

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                      TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE TITLE OF INVENTION: RELATED DISEASES FILE REFERENCE: P1982H US
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2005-04-28
FRIOR PPLICATION NUMBER: PCT/US03/34381
PRIOR PLILING DATE: 2003-10-28
PRIOR FILING DATE: 2003-10-29
PRIOR FILING DATE: 2003-10-29
NUMBER OF SEQ ID NOS: 2442
SEQ ID NO 452
LENGTH: 2000
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Pred. No. 63;
5; Mismatches 5; Indels
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Pred. No.
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APPLICANT: AGGARWAL, SUDEETA; APPLICANT: GLARK, HILARY; APPLICANT: GUNENEY, AUSTIN L.; APPLICANT: WILLIAMS, P. MICKEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1216, Application US/10533069
Publication No. US20070185017A1
GENERAL INFORMATION:
APPLICANT: AGGARWAL, SUDEEPTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88 PALENNKEQIOKIIETLTQQLQ 109
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Bést Local Similarity 44.**
Thes 8; Conservative
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Best Local Similarity 31.84
Matches 7; Conservative
WU, THOMAS D.
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; ORGANISM: Homo sapien
US-10-533-069-1216
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, ORGANISM: Homo sapien
US-10-533-069-452
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APPLICANT:
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Sequence 1242, 4pplication US/10533069

publication No. US20070185017A1

GENERAL IRPORANTION:

APPLICANT: CLARK, HILLAY

APPLICANT: CLARK, HILLAY

APPLICANT: GURNEY, AUSTIN L.

APPLICANT: GURNEY, AUSTIN L.

APPLICANT: WILLIAMS, P. MICKEY

APPLICANT: WOOD, WILLIAM I.

PRELICANT: WOOD, WILLIAM I.

PRELICANT: WOOD, WILLIAM I.

PRICE REFERENCE: P1982R1 US

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE

FRICK APPLICATION NUMBER: US/10/533,069

CURRENT APPLICATION NUMBER: US 60/422,472

PRIOR APPLICATION NUMBER: US 60/422,472

PRIOR PILING DATE: 2002-10-29

NUMBER OF SEQ ID NOS: 2442

FROMMER OF SEQ ID NOS: 2442
                                                         APPLICANT: WOOD, WILLIAM I.
APPLICANT: WU, THOMAS D.
ITITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE
ITITLE OF INVENTION: RELATED DISEASES
ITITLE OF INVENTION: RELATED DISEASES
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT APPLICATION NUMBER: PCT/US03/34381
PRIOR PILING DATE: 2003-10-28
PRIOR FILING DATE: 2003-10-29
NUMBER OF SEQ ID NOS: 2442
SEQ ID NO 1043
SEQ ID NO 1043
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Publication No. US20070185017A1
GENERAL INFORMATION:
APPLICANT: AGGARWAL, SUDEEPTA
APPLICANT: CLARK, HILARY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 42.9
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 50.0
Matches 8; Conservative
   SCHOENFELD, JILL
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ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
, ORGANISM: Homo sapien
US-10-533-069-1043
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CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2005-04-28
PRIOR FILING DATE: 2005-10-28
PRIOR FILING DATE: 2002-10-28
PRIOR FILING DATE: 2002-10-29
PRIOR FILING DATE: 2002-10-29
SEQ ID NOS: 2442
SEQ ID NO 432
                                                                                            13; Gaps
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Pred. No. 54;
4; Mismatches 3; Indels 13
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612 LLOEFRELLQYRDNSDKTLEANEMLLEKLRORI 644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 432, Application US/10533069; Publication No. US20070185017A1; GENERAL INPORMATION: APPLICANT: ACGARWAL, SUDEEPTA; APPLICANT: GLARK, HILARY; APPLICANT: GURNEY, AUSTIN L.; APPLICANT: WILLIAMS, P. MICKEY; APPLICANT: WOOD, WILLIAMS, P. MICKEY; APPLICANT: WOOD, WILLIAMS, D. APPLICANT: WOOD, WILLIAMS, D. APPLICANT: WOOD, WILLIAMS, D. APPLICANT: WILLIAMS, D. APPLICANT: WILLIAMS, D. APPLICANT: WILLIAMS, D. APPLICANT: WOOD, WILLIAMS, D. APPLICANT: WOOD, WILLIAMS, D. APPLICANT: WIL
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WILLIAMS, P. MICKEY WOOD, WILLIAM I. WU, THOMAS D.
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APPLICANT: CLARK, HILARY
APPLICANT: GURNEY, AUSTIN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 40.94
Matches 9; Conservative
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Best Local Similarity 39.4 Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
CORGANISM: Homo sapien
US-10-533-069-1070
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RESULT 8

APPLICANT: APPLICANT:

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APPLICANT: WU, THOMAS D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE
TITLE OF INVENTION: RELATED DISEASES
TITLE OF INVENTION: RELATED DISEASES
FILE REFERENCE: P1982R1 US
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT PILING DATE: 2005-04-28
FRIOR APPLICATION NUMBER: PCT/US03/34381
FRIOR APPLICATION NUMBER: US 60/422,472
FRIOR APPLICATION NUMBER: US 60/422,472
FRIOR APPLICATION NUMBER: US 60/422,472
FRIOR FILING DATE: 2002-10-29
NUMBER OF SEQ ID NOS: 2442
SEQ ID NO 565
LENGTH: 997
TYPE: PRT
CORGANISM: Homo sapien
US-10-533-069-565
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APPLICANT: GURNEY, AUSTIN L.
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: GURNEY, AUSTIN L.
APPLICANT: WOOD, WILLIAM I.
APPLICATION NUMBER: US/10/533,069
CURRENT APPLICATION NUMBER: PCT/USO3/34381
PRIOR FILING DATE: 2003-10-28
PRIOR PRILING DATE: 2002-10-29
NUMBER OF SEQ ID NOS: 2442
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            34.3%; Score 36; DB 1; Length 997;
43.8%; Pred. No. 76;
tive 3; Mismatches 6; Indels
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; Sequence 141, Application US/10533069
; Publication No. US20070185017A1
; GENERAL INFORMATION:
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                                                                     WILLIAMS, P. MICKEY WOOD, WILLIAM I.
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                       GURNEY, AUSTIN L. SCHOENFELD, JILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 43.8<sup>3</sup>
Transer 7, Conservative
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Best Local Similarity 38.5
Matches 10; Conservative
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ORGANISM: Homo sapien
US-10-533-069-141
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APPLICANT: CLARK, HILARY
APPLICANT: CHRNEY, AUSTIN 1.
APPLICANT: SCHOENFELD, JILL
APPLICANT: WOLD, WILLIAMS, P. MICKEY
APPLICANT: WOLD, WILLIAMS, P. MICKEY
APPLICANT: WO, THONAS D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE
FILE REPRENCE: P1992R1 US
FILE REPRENCE: P1992R1 US
CURRENT APPLICATION NUMBER: US/10/533,069
CURRENT FILING DATE: 2005-04-28
PRIOR PRILING DATE: 2005-028
PRIOR APPLICATION NUMBER: US 60/422,472
PRIOR FILING DATE: 2002-10-29
PRIOR FILING DATE: 2002-10-29
NUMBER OF SEQ ID NOS: 2442
SEQ ID NO 1158
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                                                                                                               TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF IMMUNE TITLE OF INVENTION: RELATED DISEASES FILE REFERENCE: P1992R1 US CURRENT APPLICATION NUMBER: US/10/533,069 CURRENT APPLICATION NUMBER: PCT/US03/34381 PRIOR FILING DATE: 2005-04-28 PRIOR FILING DATE: 2003-10-28 PRIOR FILING DATE: 2002-10-29 PRIOR FILING DATE: 2002-10-29 SEQ ID NOS: 2442 SEQ ID NO 1156 LENGTH: 491
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Pred. No. 34;
4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 36; DB 1; Length 491;
Pred. No. 34;
4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 1158, Application US/10533069; Publication No. US20070185017A1; GENERAL INFORMATION: APPLICANT: AGGARWAL, SUDEEPTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 565, Application US/10533069; Publication No. US20070185017A1
GENERAL INFORMATION: APPLICANT: APPLICANT: AGGARWAL, SUDEEPTA
                                              WILLIAMS, P. MICKEY WOOD, WILLIAM I. WU, THOMAS D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            34.3%;
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Best Local Similarity 50.0%;
Matches 6; Conservative
GURNEY, AUSTIN L. SCHOENFELD, JILL
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361 LQEIVEGMKQKM 372
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361 LQEIVEGMKQKM 372
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US-10-533-069-1158
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Best Local Similarity
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US-10-533-069-1156
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GENERAL INFORMATION

RESULT 12

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RESULT 15
US-11-657-313-24
Sequence 24, Application US/11657313
Sequence 24, Application US/11657313
Publication No. US20070186295A1
GENERAL INFORMATION:
APPLICANT: Keanrey, Paul E.
APPLICANT: Hondid, Lyes
APPLICANT: Hondy, Datrice
TILLE OF INVENTION: TAT-036 and Methods of Assessing and Treating Cancer
PILE REFERENCE: 50111/090002
CURRENT APPLICATION NUMBER: US/11/657,313
CURRENT PILING DATE: 2006-01-25
FRIOR FILING DATE: 2006-01-25
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 24
LENGRIH: 2030
TWOND THE TOWN T
APPLICANT: Chelsky, Daniel
APPLICANT: Kearney, Paul E.
APPLICANT: Paramithiotis, Eustache
APPLICANT: Paramithiotis, Eustache
APPLICANT: Hamaidi, Lyes
APPLICANT: Lamoix, Joel
APPLICANT: Lanoix, Joel
APPLICANT: Lanoix, Joel
APPLICANT: Lanoix, Joel
APPLICANT: Hugo, Parrice
TITLE OF INVENTION: TAT-036 and Methods of Assessing and Treating Cancer;
FILE REFERENCE: 5011/090002
CURRENT APPLICATION NUMBER: US/11/657,313
CURRENT APPLICATION NUMBER: US 60/762,017
PRIOR APPLICATION NUMBER: US 60/762,017
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.3
SEQ ID NO 3
LENGTH: 1359
TYPE: PRI
COGANISM: Homo sapiens
US-11-657-313-3
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LOCATION: (896)..(896)
OTHER INFORMATION: Xaa can be any naturally occurring amino acid
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33.8%; Score 35.5; DB 7; Length 2030;
Best Local Similarity 52.4%; Pred. No. 2e+02;
Matches 11; Conservative 6; Mismatches 1; Indels 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 33.8%; Score 35.5; DB 7; Length 1359; Best Local Similarity 52.4%; Pred. No. 1.3e+02; Matches 11; Conservative 6; Mismatches 1; Indels 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1278 ILDL--DLLLELLE-LQERLR 1295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              : | | | : | | | | | | : : | : | | | 435 ILDL--DLLLELLE-LQERLR 452
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Search completed: August 16, 2007, 22:55:51 Job time : 7 secs

# Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- permanent accession numbers. The new UniProtrecord may not contain the previous temporary numbers from the most recent version of UniProt. These sequences have been assigned new Please note that the curators of the UniProt database have purged some temporary accession
- extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein If you encounter an accession number from an older search run against UniProt (results file Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

members of the public who may encounter UniProt temporary accession When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or numbers

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GO; GO:0003723; F:RNA binding; IEA.
INLECTPO; IPR003890; IP_EIF4G.
Pfam; PF02854; MIF4G; 2.
SMART; SM00543; MIF4G; 1.
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Q1VGCO_V1BAL
Q87FD0_V1BPA
Q1AUZ7_9ACTN
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Ehrlich S.D., Berka R.M.;
"Complete genome sequence of the industrial bacterium Bacillus
licheniformis and comparisons with closely related Bacillus species.";
Genome Biol. S:RESEARCH077.1-RESEARCH077.12(2004).
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PubMed=15461803; DOI=10.1186/gb-2004-5-10-r77;
Rey MW., Ramaiya P., Nelson B.A., Brody-Karpin S.D., Zaretsky E.J., Tang M., Lopez de Leon A., Xiang H., Gusti V., Clausen I.G., Olsen P.B., Rasmussen M.D., Andersen J.T., Joergensen P.L.,
                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina Hammon N., Israni S., Pitluck S., Saunders E.H., Schmutz J., Latimer F., Land M., Kyrpides N., Mavrommatis K., Richardson P.; "Complete sequence of Anabaena variabilis ATCC 29413.";

*Submitted (SEP-2005) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                      Anabaena variabilis (Etrain ATCC 29413 / PCC 7937).
Bacteria: Cyanobacteria: Nostocales; Nostocaceae; Anabaena.
NCBL_TaxID=240292;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=15383718; DOI=10.1159/000079829;
Veith B., Herzberg C., Steckel S., Feesche J., Maurer K.H.,
Ehrenzeich P., Baeumer S., Henne A., Liesegang H., Merkl R.
Ehrenzeich A., Gottschalk G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 892;
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Bacillus licheniformis (strain DSM 13 / ATCC 14580).
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Complete protecome; Hypothetical protein.
SEQUENCE 892 AA; 104914 MW; 456F1AD7C0451004 CRC64;
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                                                                                                               25-OCT-2005, integrated into UniProtKB/TrEMBL
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                                                                    PRT; 892 AA.
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                                                                                                                                            , sequence version 1.
                                                                  PRELIMINARY;
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                                                                                                                                                          02-MAY-2006, entry version 6.
Hypothetical protein.
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QESPC9; Q62ZR7;
                                                             Q3MCF8 ANAVT
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                                                             Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
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Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Buarchontoglires, Glires, Rodentia, Sciurognathi,
Muroidea, Muridae, Murinae, Rattus.
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MEDINE=98077648; PubMed=9415807;
Chiang A.N., Fan K.C., Shaw G.C., Yang U.C.;
"Repetitive elements in the third intron of murine apolipoprotein A-I
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                                                                                                                                                       EMBL; AE017333; AAU39085.1; -; Genomic_DNA.

REMBL; CP000002; AAU21741.1; -; Genomic_DNA.

RenomeReviews; CP000002 GR; BL03268.

RenomeReviews; AE01733 GR; BL100111.

RG0; G0:0005737; C::qvcpplasm; IEA.

RG0; G0:0016740; F::rerine O-acetyltransferase activity; IEA.

RG0; G0:0016740; F::ransferase activity; IEA.

RG0; G0:0016515; P::qysteine biosynthesis from serine; IEA.

InterPro; IPR001801; CysE transferase.

R InterPro; IPR001801; Hexapep_transf.

R InterPro; IPR001004; Trimer_LpxA_like.

PFam: PR00132; Hexapen-. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51.9%; Score 54.5; DB 2; Length 216; 59.1%; Pred. No. 61; ive 4; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00132; Hexapep; 4. ____TIGREAMs; TIGR01172; cysE; 1.
Acyltransferase; Amino-acid biosynthesis; Complete proteome;
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STRAIN-WKY, and SHRSP, TISSUE-Spleen;
Chiang A.-N., Fan K.-C., Shaw G.-C., Yang U.-C.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cysteine biosynthesis; Repeat; Transferase.
SEQUENCE 216 AA; 24124 MW; 4C1ADOCC51F74B4C CRC64;
-!- SUBCELLULAR LOCATION: Cytoplasm (By similarity).
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GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0008289; F:lipid binding; IEA.
GO; GO:0006869; F:lipid transport; IEA.
GO; GO:0042157; P:lipoprotein metabolism; IEA.
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                                                                                                                                             258 AA.
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EMBL; U79577; AAB58429.1; -; Genomic_DNA.
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InterPro; IPR000074; ApoAl A4 E.
Pfam; PF01442; Apolipoprotein; 1.
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                                     SEQUENCE
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                   NCBI_TaxID=10116;
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SIGNAL
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Distributed under the Creative Commons Attribution-Noberivs License
                                                                                                                                                                               Rattus norvegicus (Rat).
Bukaryota, Metazoa; Chordata, Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                    STRAIN=SHR; TISSUE-Spleen;
MEDLINE=98077648; PubMed=9415807;
Chiang A.N., Fan K.C., Shaw G.C., Yang U.C.;
"Repetitive elements in the third intron of murine apolipoprotein A-I
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
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                    Length 258;
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54.5%; Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                             STRAIN=SRR; TISSUE=Spleén;
Chiang A.-N., Fan K.-C., Shaw G.-C., Yang U.-C.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               258 AA; 29831 MW; 093FB582E629CDC8 CRC64;
29918 MW; 093E6EF2E629CDC8 CRC64;
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                  50.5%; Score 53; DB 2; Ler 54.5%; Pred. No. 1.1e+02; tive 4; Mismatches 6;
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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I)
                                                                                                                                                                                                                                                                                                                                                                                                                 GO, GO:0005576; C:extracellular region; IEA.
GO; GO:0008289; F:lipid binding; IEA.
GO; GO:0006669; P:lipid transport; IEA.
GO; GO:0042157; P:lipoprotein metabolism; IEA.
InterPro; IPR013126; ApoAl E ApoLp.
InterPro; IPR000074; ApoAl Ā4 E.
                                                                                                                                           01-JUL-1997, integrated into UniProtKB/TrEMBL. 01-JUL-1997, sequence version 1.
                                                                                                                           258 AA.
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HSSP; P02647; 1AV1.
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121 PHLDEFQEKWNEEVEAYRQKLE 142
                                                                                                                           PRELIMINARY; PRT;
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                                                        1 PVLDLFRELLNELLEALKQKLK 22
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                                       12; Conservative
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 258 AA;
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Best Local Simi
Matches 12;
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P04639;
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Pfam; PF0142; Apolipoprotein; 1.
Cfolesterol metabolism; Direct protein sequencing; HDL;
Lipid metabolism; Lipid transport; Repeat; Signal; Steroid metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The primary translation product of rat intestinal apolipoprotein A-I mRNA is an unusual preproprotein.";
J. Biol. Chem. 25:971-978 (1982).
-!-FUNCTION: Participates in the reverse transport of cholesterol from tissues to the liver for excretion by promoting cholesterol efflux from tissues and by acting as a cofactor for the lecithin cholesterol acyltransferase (LCAT).
-!-SUBCELIULAR LOCATION: Secreted protein.
-!-TISSUE SPECIFICITY: Major protein of plasma HDL, also found in
                                                                                                                                                                            MEDLINE-84207987; PubMed-6426956;
Poncin J.E., Martial J.A., Gielen J.E.;
"Cloning and structure analysis of the rat apolipoprotein A-I CDNA.";
Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
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                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLINE=87008540; PubMed=3020028;
Haddad I.A., Ordovas J.M., Pitzpatrick T., Karathanasis S.K.;
"Linkage, evolution, and expression of the rat apolipoprotein A-I, III, and A-IV genes.";
J. Biol. Chem. 261:13268-13277(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- SIMILARITY: Belongs to the apolipoprotein Al/A4/E family.
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EMBL; X00558; CAA25224.1; -; mRNA.

EMBL; X0259; AAA40745.1; -; Genomic_DNA.

PIR; A24700; A24700.

HSSP; P02647; 1AV1.

RGD; GO: 0017127; F: cholesterol transporter activity; IDA.

GO; GO: 0005548; F: phospholipid transporter activity; IDA.

GO; GO: 0015914; P: phospholipid transporter activity; IDA.

GO; GO: 0015914; P: phospholipid transport; IDA.

RGO; GO: 0015914; P: phospholipid transport; IDA.

RGO; GO: 0015914; P: phospholipid transport; IDA.

RIMEEPPO; IPRO11326; ApoA/E ApoLp.

RIMEEPPO; IPRO100074; ApoAl A4 E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=Sprague-Dawley;
MEDINE=82098162; PubMed=6798036;
Gordon J I., Smith D.P., Andy R., Alpers D.H., Schonfeld G.,
Strauss A.W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -> K (in Ref. 2).
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-> K (in Ref. 2).
2E8D5EB45FEAEE88 CRC64;
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Apolipoprotein A-I.
/FTId=PRO_000001954.
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                                                                                                                                                                                                                                                                                           Eur. J. Biochem. 140:493-498(1984)
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122 PHLDEFQEKWNEEVEAYRQKLE 143
1 PVLDLFRELLNELLEALKOKLK 22
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SEQUENCE
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Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Altschul S.F., Zeeberg B. Buetow K.H., Schaefer C.F., Bata N.K.,
Altschul S.F., Zeeberg B. Buetow K.H., Schaefer C.F., Bata N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haidh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Brownstein M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toohiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Rockers R.W., McKernan K.J., Malek J.A., Gunzatne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Richards S., Worley K.C., Hale S., Garchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Brown A., Schein J.B., Jones S.J.M., Marra M.A.,
Brown A., Schein J.B., Jones E.J.M., Marra M.A.,
Brown A., Schein J., Jones E.J.M., Marra M.A., Schein J., Jones E.J.M., Marra M.A.,
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Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus norvegicus (Rat).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
Muroidea, Muridae, Murinae, Rattus.
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                                50.5%; Score 53; DB 1; Length 259; 54.5%; Pred. No. 1.1e+02; tive 4; Mismatches 6; Indels
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54.5%; Pred. No. 1.18+02;
ive 4; Mismatches 6; Indels
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Director MGC Project;
Submitted (FEB-2005) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        259 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRELIMINARY; PRT;
                                                                                                                                                                                                                                              | || |:| || :|| :|| || PHLDEFQEKWNEEVEAYROKLE 143
                                                                                                                                                                                             1 PVLDLFRELLNELLEALKQKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR013326; ApoA/E_ApoLp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry version 6.
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                                                                     Best Local Similarity 54.5
Matches 12; Conservative
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Les 12; Conserv
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SEQUENCE 2
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MEDLINE=21537279; PubMed=11679669; DOI=10.1126/science.1063447;

A Glaser P., Frangeul L., Buchrieser C., Rueniok C., Amend A.,

Baquero F., Berche P., Blocker H., Brandt P., Chakraborty T.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Brian K.-D., Fshi H., Garcia-Clopez N., Hain T., Hauf J., Jackson D.,

Gautier L., Goobel W., Gomez-Lopez N., Hain T., Hauf J., Jackson D.,

Andueno E., Mattournam A., Mata Vicente J., Ng E., Nedjari H.,

Nordaick G., Novella S., de Pallos B., Perez-Diaz J.-C., Purcell R.,

Remmel B., Rose M., Schlueter T., Simoes N., Tierrez A.,

Remmel B., Rose M., Schlueter T., Simoes N., Tierrez A.,

Raquez-Boland J.-A., Voss H., Wehland J., Cossart P.,

"Comparative genomics of Listeria species.";

Science 294:849-852(2001).

C. -1- SIMILARITY: Belongs to the UPF0085 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Listeria monocytogenes serotype 4b (strain F2365).
Bacteria, Firmicutes, Bacillales, Listeriaceae, Listeria.
                                                                                                                                                                                                                                                                Bacteria; Firmicutes; Bacillales; Listeriaceae; Listeria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         149 156 ATP (Potential).
274 AA; 30436 MW; A149AF0F8D5D1B1A CRC64;
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Y1457 LISMO STANDARD; PRT; 274 AA.
P67195; 092BQ3;
11-OCT-2004, integrated into UniProtKB/Swiss-Prot.
11-OCT-2004, sequence version 1.
30-MAY-2006, entry version 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-2004, integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
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ListiList; LMO01457; -.
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07-MAR-2006, entry version 15.
UPF0085 protein LMOE2365 1476.
OrderedLocusNames-LMO62365 1476;
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InterPro; IPR005177; DUF299.
Pfam; PF03618; DUF299; 1.
                                                                                                                                                                                                      OrderedLocusNames=1mo1457;
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                                                                                                                                                                 UPF0085 protein lmo1457
                                                                                                                                                                                                                                        Listeria monocytogenes.
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Best Local Similarity
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274 AA; 30436 MW; A149AP0F8D5D1B1A CRC64;
                                                                                                                                                                                                                                                                                 UPF0085 protein lin1494.
/FTId=PRO_0000196672.
                                                                                                       EMBL; AL596168; CAC96725.1; -; Genomic_DNA
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Best Local Similarity 55.0°
Matches 11; Conservative
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                                                                                   PIR; AE1619; AE1619
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MEDLINE=21537279; PubMed=11679669; DOI=10.1126/science.1063447;
Glaser P., Frangeul L., Buchrieser C., Rusniok C., Amend A.,

Baquero F., Berche P., Blocker H., Brandt P., Chakraborty T.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Charbit A., Chetouani F., Couve E., de Daruvar A., Dehoux P.,

Butian R., Dominguez-Bernal G., Duchaud E., Durant L., Dussurget O.,

A Gautiar L., Goebel W., Ganzcia-Copez N., Hain T., Hauf J., Jackson D.,

A Jones L.-M., Kaerst U., Kreft J., Kuhn M., Kunst F., Kurapkat G.,

A Madueno E., Maitournam A., Mata Vicente J., Ngg E., Nedjari H.,

Nordsiek G., Novella S., de Pablos B., Perez-Diaz J.-C., Purcell R.,

Remmel B., Rose M., Schlueter T., Simoes N., Tierrez A.,

A Vazquez-Boland J.-A., Voss H., Wehland J., Cossart P.,

"Comparative genomics of Listeria species.";

I. Science 294:849-852 (2001).

Corpusarative Service Comparative Comparativ
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                                                                                PubMed=15118801; DOI=10.1093/nar/gkh562;
Nelson K.E., Fouts D.E., Mongodin B.F., Ravel J., DeBoy R.T.,
Kolonay J.F., Rasko D.A., Angluoli S.V., Gill S.R., Paulsen I.T.,
Brinkac L.M., Daugherty S.C., Dodson R.J., Durkin A.S., Madupu R.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Forberger H.A., Tran B., Kathariou S., Wonderling L.D., Uhlich G.A.,
Bayles D.O., Luchansky J.B., Fraser C.M.;
"Whole genome comparisons of serotype 4b and 1/2a strains of the food
borne pathogen Listeria monocytogenes reveal new insights into the
core genome components of this species.";
Nucleic Acids Res. 32:2386-2395(2004).
-! SIMILARITY: Belongs to the UPF0085 family.
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11-OCT-2004, sequence version 1.
07-M8K-2006, entry version 7.
UPF0085 protein lin1494.
                                                               NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AE017262; AAT04251.1; -; Genomic_DNA.
GenomeReviews; AE017262_GR; LMOF2365_1476.
TIGR; LMOF2365_1476; -.
HAMAP; MF 01062; -; 1.
InterPro; IPR005177; DUF299.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OrderedLocusNames=lin1494;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
     NCBI_TaxID=265669;
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Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; DOI=10.1093/nar/gkh562;
Whomed-15115801; Doi-10.1093/nar/gkh562;
White O., Nelson W.C., Nierman W.C., Beanan M.J.,
Brinkac L.M., Daudherty S.C., Dodson R.C., Madupu R.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., The Sequence shown here is derived from an EmblyGensan K.E.

EmblyGensan K.D., June M. L., Man S.E., Man J., June J.
EmblyGensan K.D., June J., 
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Distributed under the Creative Commons Attribution-NoDerivs License
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Score 53; DB 1; Length 274;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Listeria monocytogenes str. 4b H7858.
Bacteria; Firmicutes; Bacillales; Listeriaceae; Listeria.
NCBI_TaxID=267410;
                                                                                                                   6; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-SEP-2005, integrated into UniprotKB/TrEMBL. 13-SEP-2005, sequence version 1. 18-APR-2006, entry version 3. Hypothetical protein.
                                                                                                                   3; Mismatches
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Q4EGJ5;
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EMBL; CP000094; ABA75953.1; -; Genomic_DNA.
GenomeReviews; CP000094 GR; Pf1 4216.
InterPro; IPR011029; DEATH_like.
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                                                                                        InterPro; IPR012199; MazG.—
InterPro; IPR011551; MazG bac.
InterPro; IPR004518; MazG_cat.
                                                                                                                                                                                                                                                                   280 AA; 32085 MW;
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Q9F2I5 arath
ID Q9F2I5_arath preliminary;
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                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                              PERMINEL/2a F6854;
PubMed=15115801; DOI=10.1093/nar/gkh562;
PubMed=15115801; DOI=10.1093/nar/gkh562;
PubMed=15115801; DOI=10.1093/nar/gkh562;
Rolson K.E., Route D.E., Morgodin E.F., Ravel J., DeBoy R.T.,
Kolonay J.F., Rasko D.A., Angiuoli S.V., Gill S.R., Paulsen I.T.,
Peterson J.D., White O., Nelson W.C., Nierman W.C., Beanan M.J.,
Brinker L.M., Daugherty S.C., Dodson R.J., Durkin A.S., Madupu R.,
Haft D.H., Selengut J., Van Aken S.E., Khouri H.M., Fedorova N.,
Forberger H.A., Tran B., Karhariou S., Wonderling L.D., Uhlich G.A.,
Bayles D.O., Luchansky J.B., Fraser C.M.;
"Whole genome comparisons of sercetype 4b and 1/2a strains of the food-
borne pathogen Listeria monocytogenes reveal new insights into the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US DOE Joint Genome Institute;
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina T.,
Hammon N., Israni S., Pitluck S., Saunders E.H., Schmutz J.,
Larimer F., Land M., Kyripides N., Anderson I., Richardson P.;
"Complete sequence of Pseudomonas fluorescens Pf0-1.";
Submitted (AUG-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=Pfl 4216;
Pseudomonas fluorescens (strain PfO-1).
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
Pseudomonadaceae; Pseudomonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
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                                                                                           13-SEP-2005, integrated into UniProtKB/TrEMBL.
18-ASR-2006, sequence version 1.
18-APR-2006, entry version 3.
Hypothetical protein.
08/FNames-EMO66854 1500;
Listeria monocytogenes str. 1/2a F6854.
Bacteria, Pirmicutes; Bacillales; Listeriaceae; Listeria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Indels
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274 AA; 30436 MW; A149AF0F8D5D1B1A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50.5%; Score 53; DB 2; 155.0%; Pred. No. 1.2e+02;
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08-NOV-2005, sequence version 1.
30-MAY-2006, entry version 8.
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                                                   274 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    core genome components of this species.";
Nucleic Acids Res. 32:2386-2395(2004).
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                                          LISMO PRELIMINARY;
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Pfam; PF03618; DUF299; 1.
Hypothetical protein.
SEQUENCE 274 AA; 30436 MA
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Best Local Similarity 55.04
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                   NCBI_TaxID=267409;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q3K8FL PSEPF
ID G3K8FL PSEPF
DT 08-NOV-2005, in'
DT 08-NOV-2005, se-
DT 08-NOV-2006, en 
DE MaZG.
GN OrderedLocusNam
OC Bacteria, Prote
OC Pseudomonas flu
OC Pseudom
                             LISMO
                                            Q4ETW3
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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (FEB-2006) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                                           50.5%; Score 53; DB 2; Length 280; 57.9%; Pred. No. 1.2e+02; ive 4; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 605;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Siefert J., Ferriera S., Johnson J., Kravitz S., Halpern
Remington K., Beeson K., Tran B., Rogers Y.-H., Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50.5%; Score 53; DB 2; Length 605
57.1%; Pred. No. 2.5e+02;
ive 2; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUN-2006, entry version 3.
Oligoendopeptidase F.
ORRNames=B14911_28130;
Bacillus sp. NRRL B-14911.
Bacteria, Firmicutes; Bacillales; Bacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GO, GO:0004222; F:metalloendopeptidase activity; IEA.
GO; GO:0006508; P:proteolysis; IEA.
InterPro; IPR001667; Pept MAA M3B.
InterPro; IPR018647; Peptidase M3 N.
InterPro; IPR018647; Peptidase M3 N.
InterPro; IPR004438; Peptidase M3 N.
Pfam; PF01422; Peptidase M3; 1.
Pfam; PF01422; Peptidase M3; 1.
ITGRFAMS; ITGR00181; Peptidase M3; 1.
ITGRFAMS; ITGR00181; Peptidase M3 N. 1.
ITGRFAMS; ITGR00181; Peptidase M3 N. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           300142; ZINC PROTEASE; UNKNOWN 1.
605 AA; 68388 MW; 36DEA70199190077 CRC64;
                                                                                      0AA804512A53414D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AAOX01000006; EAR66951.1; -; Genomic DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-APR-2006, integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      727 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   605 AA.
PIRSF; PIRSF002844; NTP_pyrphdr_MazG; 1.
TIGRFAMs; TIGR00444; mazG; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QZBAJ7 9BACI PRELIMINARY; PRT; Q2BAJ7;
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Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                         Recensories N.A., Palm C.J., Conway A.B., Conn L., Hansen N.F., Altafi H., Nguyen M., Lam B., Southwick A., Miranda M., Brooks S., Buehler E., Chao Q., Chin C., Chiou J., Choi B., Gonzalez A., Howng B., Johnson-Hopson C., Khan S., Kim C., Koo T., Lee J.M., Lenz C., Liu A., Liu S., Mukharsky N., Pham P., Sakano H., Shinn P., Toriumi M., Vaysberg M., Yu G., Ecker J., Theologis A., Davis R.W.; Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R GO; GO: 00005524; F.APP binding; IEA.

R GO; GO: 00005524; F.APP binding; IEA.

R GO; GO: 0000166; F: Protein serine(hreonine kinase activity; IEA.

R GO; GO: 00064674; F: Protein serine(hreonine kinase activity; IEA.

R GO; GO: 0006468; P: Protein amino acid phosphorylation; IEA.

R GO; GO: 0006468; P: Protein amino acid phosphorylation; IEA.

R InterPro; IPR00009; Kinase like.

R InterPro; IPR00019; Protein Ans.

R Prodom; P0000001; Protein Kinase; 1.

R PROSITE; PS00101; PROTEIN KINASE ATP; 1.

R PROSITE; PS00101; PROTEIN KINASE DOM; 1.

R PROSITE; PS00108; PROTEIN KINASE DOM; 1.

R PROSITE; PS00108; PROTEIN KINASE ST; 1.

R PROSITE; PS00108; PROTEIN KINASE ST; 1.

R ATP-binding; Hypothetical protein; Kinase; Nucleotide-binding; Serine/threonine-protein kinase; Transferase.
                                                                                                                                             Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicotyledons,
rosids, eurosids II, Brassicales, Brassicaceae, Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50.5%; Score 53; DB 2; Length 727; 52.4%; Pred. No. 3e+02; tive 4; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              727 AA; 81129 MW; BEF55B9646E0459E CRC64;
Q9FZI5;
01-MAR-2001, integrated into UniProtKB/TrEMBL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AC007152; AAF98207.1; -; Genomic_DNA. HSSP; P06239; 3LCK.
                                        01-MAR-2001, sequence version 1.
18-APR-2006, entry version 22.
Hypothetical protein F1019.1 (Fragment)
                                                                                                       Name=F1019.1;
Arabidopsis thaliana (Mouse-ear cress)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 52.44
                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                      NCBI_TaxID=3702;
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239 PLLDRFRGVLNNLREMCKRKV 259 셤

1 PVLDLFRELLNELLEALKOKL 21

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Gaps

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Search completed: August 16, 2007, 22:48:32 Job time : 351 secs

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nCore version	- 2007
GenCore	(c) 1993
	Copyright (

OM protein - protein search, using sw model

Run on:

August 16, 2007, 22:38:56 ; Search time 213 Seconds (without alignments) 50.541 Million cell updates/sec

US-10-715-895A-4 Title: Perfect score:

105 1 PVLDLPRELLNELLEALKOKLK 22 Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

2782304 seqs, 489333398 residues Searched:

2782304 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

A\_Geneseq\_200701:\* geneseqp1980s:\* geneseqp1990s:\* •• Database

geneseqp2006s:\* geneseqp2003bs:\* geneseqp2003as:\* geneseqp2000s:\* geneseqp2001s:\* geneseqp2002s:\* geneseqp2004s:\* geneseqp2005s:\* .... 11:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

SUPERSTERS	Descript on		Aay01708		_	AAY18779 Lecithin	AAY18963 Lecithin	AAY18937 Lecithin	AAY19033 Lecithin	AAY19191 Lecithin	AAY19217 Lecithin	AAY19287 Lecithin	AAY18446 Lecithin	AAY18516 Lecithin		ADG20900 Apolipopa			ADJ32938 Apo lipop	Apo	Apo	Apo	Apo	
ני	£	ì	AAYO	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	AAY1	ADG2	ADG	ADG	ADJ3	ADJ	ADJ:	A C	ADG	
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	& Query Match Length DB	To Grand	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	22	
•	Query	Harch	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	
	1 0 0	1000	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	105	
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	22222
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2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	4 4 4 4 4 0 11 12 12 14 12

### ALIGNMENTS

AAY01708 standard; peptide; 22 AA (first entry) 24-JUN-1999 AAY01708; RESULT 1 AAY01708 

Peptide for making peptide-lipid complex by co-lyophilization approach

Peptide-lipid complex; co-lyophilization approach; liposome; compound strorage; vaccine; dyslipoproteinemia; hypercholesterrolemia; phypertriglyceridemia; az low HDL; apolipoprotein A-1 deficiency; cardiovascular disease; atherosclerosis; septic shock; infectious disease.

Synthetic.

WO9917740-A1

98WO-US020330. 28-SEP-1998; 15-APR-1999.

97US-00942597. 02-OCT-1997;

(DASS/) DASSEUX J.

Dasseux J;

WPI; 1999-277181/23.

Preparation of a lyophilized peptide/lipid product - by co-lyophilization of peptides and solubilizing in lipid.

Example 1; Page 18; 42pp; English.

The present sequence represents a peptide used to make peptide-lipid complexes by the co-lyophilization approach of the invention. Preparation of a lyophilized peptide/lipid product comprises co-lyophilization of one or more peptides which are able to adopt an amiphipathic conformation or their analogues, and one or more lipids in a solvent system to form a peptide/lipid product which can be rehydrated to form peptide/lipid product which can be rehydrated to form peptide/lipid complexes, solubilizing at least one amphipathic peptide or its analog in a first solution, solubilizing at least one lipid in a second solution

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Pyophilizing to form a product which can be rehydrated to form peptide/lipid complexes. The method is used for for generating stable peptide/lipid vesicles and complexes such as micellar, spherical and discoidal complexes in bulk preparations and insmaller units e.g. dosages. Liposomes are known to be used for delivery vehicles for drugs, cosmetics and bloactive compounds. The method may also be used for accommetics and bloactive compounds. The method may also be used for provantion of lipids or for formulation of products for treatment or prevention of human diseases such as co-presentation of antigens in vaccines, treatment or prevention of fysiloporceinnalas, e.g. typercholesterolemia, hypertriglyceridemia, low HDL and apoliporocein A-1 deficiency, cardiovascular disease such as atherosclerosis, septic shock or infectious diseases. The method can also be used in the preparation of complexes which can be used as carriers for drugs, as vectors (to deliver drugs, DNA and genes) e.g. to the liver or to extra hepatic cells or as scavengers to trap toxins such as pesticides and LPS
   combining the solutions and
   which is miscible with the first solution,
856666666666666668888
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Sequence 22 AA;

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  Length 22;
                        Indels
100.0%; Score 105; DB 2;
100.0%; Pred. No. 9.2e-07;
tive 0; Mismatches 0;
                                                22
                                                           1 PVLDLFRELLNELLEALKQKLK 22
                                               1 PVLDLFRELLNELLEALKOKLK
                       22; Conservative
           Similarity
Query Match
Best Local
                       Matches
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Gaps

RESULT 2

AAY18683 standard; peptide; 22 AA

AAY18683;

Lecithin: cholesterol acyltransferase activation exhibiting peptide #4. (first entry) 09-JUL-1999

Apolipoprotein A-1; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; UCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDi; apoA-1; high density lipoprotein; hypertriglyceridemia; metabolic syndrome; septic shock 

Synthetic

Homo sapiens

W09916408-A2

98WO-US020328 28-SEP-1998; 97US-00940093 29-SEP-1997;

DASSEUX J. (DASS/) (SEKU/) (BUTT/)

SEKUL R. BUTTNER K. CORNUT I. METZ G. (METZ/) (CORN/)

Cornut I, Buttner K, Sekul R, Dasseux J,

ΰ

WPI; 1999-277031/23.

Peptide agonists of apolipoprotein A-I.

Example; Page 104; 152pp; English.

The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 14-22 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid

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dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, resenonsis, HDL (high density lipoprotein) or appA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to inage HDL at e.g. atherosclerotic streak, and to raise antibodies. AAY18680 to AAY189313 represent lecithin.cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (apoA-1) which is a 14-22 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-1 deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used disquostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atheroscoleroois; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lecithin: cholesterol acyltransferase activation exhibiting peptide #30
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                                                                                                                                                                                                                              100.0%; Score 105; DB 2;
100.0%; Pred. No. 9.2e-07;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide agonists of apolipoprotein A-I.
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                                                                                                                                                                                                                                                                                                                                                                                                                            AAY18709 standard; peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Buttner K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98WO-US020328.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-277031/23
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BUTTNER K.
CORNUT I.
                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                               Sequence 22 AA;
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Local Sim-
22; C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           septic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9916408-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY18709;
                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (METZ/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORN/)
                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                             RESULT 3
                                                                                                                                                                                                                                                                                                                                                                                                            AAY18709
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Gaps

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Indels

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Pred. No. 9.2e-07;

10.001 ; P.S.

22; Conservative

Mismatches

22

1 PVLDLFRELLNELLEALKOKLK 22

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Best Local Similarity
Matches 22; Conserv
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AAY18963
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                                                                                                                                                                                                                                                                                 Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                            Lecithin:cholesterol acyltransferase activation exhibiting peptide #100.
 e.g. atherosclerotic streaks, and to raise antibodies. AAY18680 to AAY18933 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists
                                                                                             Gaps
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                                                                    Length 22;
                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Metz G;
                                                                    Score 105; DB 2;
Pred. No. 9.2e-07;
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                                                                                         0; Mismatches
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                                                                                                                1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                                                                                                             AAY18779 standard, peptide; 22 AA
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                                                                    100.0%;
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                                                                               Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEKUL R.
BUTTNER K.
CORNUT I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-277031/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DASSEUX J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 AA;
                                                Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                septic shock
                                                                                                                                                                                                                                       09-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jasseux J,
                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                   AAY18779;
                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CORN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BUTT/)
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100.0%; Score 105; DB 2; Length 22;

Query Match

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                                                                                                                                                                                                       Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to AAY19187 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists
                                                                                                                                                           Lecithin:cholesterol acyltransferase activation exhibiting peptide #30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes an agonist (A) of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide agonists of apolipoprotein A-I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sxample; Page 108; 254pp; English.
AAY18963 standard; peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Buttner K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00940096.
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                                                                                                          (first entry)
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BUTTNER K.
CORNUT I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             METZ G.
                                                                                                                                                                                                                                                                                                                       septic shock
                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                          09-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                      AAY18963;
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(CORN/)
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Matches
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AAY18937;

RESULT 6 AAY18937

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The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Apolipoprotein A-I; agoniat; dyalipidemic disorder; dyalipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I;
                                                                         Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                      Lecithin:cholesterol acyltransferase activation exhibiting peptide #100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to
AAY19187 represent lecithin:cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lecithin:cholesterol acyltransferase activation exhibiting peptide #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exhibiting core peptides, which are apoA-I agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 105; DB 2;
100.0%; Pred. No. 9.2e-07;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Buttner K, Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide agonists of apolipoprotein A-I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PVLDLFRELLNELLEALKOKLK 22
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                                                                                                                                                                                                                                                                                                                                                       98WO-US020326.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dasseux J, Sekul R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-277034/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEKUL R.
BUTTNER K.
CORNUT I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (METZ/) METZ G.
                                                                                                                                                                                                                                                                                                                                                       28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-1997;
                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                       WO9916458-A1
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                                                                                                                                                             septic shock
                                                                                                                                                                                                                                                                                                              08-APR-1999
                                                                                                                                                                                                         Synthetic.
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                     (DASS/)
(SEKU/)
(BUTT/)
(CORN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertrigityceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at
                                                                                                                                                                                                                      Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome; septic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to AAY19187 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists
                                                                                                                                                                                   Lecithin: cholesterol acyltransferase activation exhibiting peptide #4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes an agonist (A) of apolipoprotein A-I (apob-1) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with
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                                                            AAY18937 standard; peptide; 22 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US020326
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                                                                                                                                            (first entry)
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BUTTNER K.
CORNUT I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DASSEUX J.
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                                                                                                                                          09-JUL-1999
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(CORN/)

(DASS/) (SEKU/) (BUTT/)

Metz G;

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Gaps ; 0

Query Match

AAY19033

AAY19033 ID AAY1 XX AC AAY7 XX

RESULT 7

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Length 22; Indels Dufourcq J;

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The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to reise antibodies. AAY19188 to AAX19441 represent lecithin; cholesterol acyltransferase (LCAT) activity
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                                                                                                                                                                                                                                                                                                    Cornut I, Metz G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exhibiting core peptides, which are apoA-I agonists
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100.0%; Pred. No. 9.2e-07;
Live 0; Mismatches 0;
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                                                                                                                                                                                                BUTTNER K. CORNUT I.
                                                                                                                                                                                                                                        METZ G.
DUFOURCQ J.
                                                                                                                                                             DASSEUX J.
                                                                                                                                                                                 SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 AA;
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                                                                               28-SEP-1998;
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  WO9916459-A1
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                                          08-APR-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY19287;
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                                                                                                                                                                                                                                      (METZ/)
(DUFO/)
                                                                                                                                                           (DASS/)
(SEKU/)
                                                                                                                                                                                                                      CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY19287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAX19188 to AAX19441 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; decithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome; septic shock.
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high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                        Dufourcq J;
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100.0%; Pred. No. 9.2e-07;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide agonists of apolipoprotein A-I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 19; Page 155; 280pp; English
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                                                                                                                                                                                                    98WO-US020327.
                                                                                                                                                                                                                                           97US-00940095.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-277035/23
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Best Local Similarity
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                                                                                                                                                                                                                                                                                 DASSEUX J.
                                                                                                                                                                                                                                                                                                                      BUTTNER K.
                                                                                                                                                                                                                                                                                                                                         CORNUT 1.
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                                                                                                                                                                                                                                                                                                                                                               METZ G.
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                                                                                                                                                                                                                                           29-SEP-1997;
                                                                               Homo sapiens
                                                                                                                                                                                                    28-SEP-1998;
                                                                                                                      WO9916459-A1
                      septic shock
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                                                             Synthetic.
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                                                                                                                                                                                                                                                                                 DASS/)
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RESULT 9 AAY19217

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Gaps

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Length 22; Indels Dufourcd J;

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The present invention describes a nucleic acid (A) encoding an apolipoprotein A-I (apoA-I) agonist (B) that is a peptide, or analog, which forms an amphipathic alpha-helix in presence of lipids. (A), optionally as a complex with lipids, and host cells that contain (A), are useful for gene therapy, or prevention, of diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) and apoA-I deficiency, hypertriglyceridemia and metabolic syndrome, also to treat endotoxemia (septic shock). Host cells containing (A) can also be used to study the role of apoA-I in lipid metabolism. (B) can be used to diagnostically, e.g. to measure serum HDL (particularly its subpopulation involved in retrograde cholesterol transport) and for imaging the circulatory system or HDL accumulations at fatty streaks. The present
                         Cornut I, Metz G,
                                                                                                                Nucleic acid encoding apoproteinA-I agonist peptides
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                                                                                                                                                            English.
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                         Buttner K,
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                                                                                                                                                            Example; Page 135; 232pp;
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                         Dasseux J, Sekul R,
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                                                                   WPI; 1999-254921/21
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BUTINER K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORNUT 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 AA;
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(SEKU/)
(BUTT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ch 100.0%; Score 105; DB 2; Length 22; 1 Similarity 100.0%; Pred. No. 9.2e-07; 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                         Metz G,
                                                                                                                                                                                                      Cornut I,
                                                                                                                                                                                                                                                                                               Peptide agonists of apolipoprotein A-I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY18446 standard; peptide; 22 AA
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29-SEP-1997; 97US-00940095
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                                                                                                                                                                                                      Dasseux J, Sekul R,
                                                                                                                                                                                                                                                   WPI; 1999-277035/23
                                                                 SEKUL R.
BUTTNER K.
CORNUT I.
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BUTTNER K.
CORNUT I.
METZ G.
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                                                                                                                                                          DUFOURCQ J
                                            DASSEUX J.
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                                                                                                                                       METZ G.
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                                                                                      (BUTT/)
(CORN/)
(METZ/)
(DUFO/)
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                                                             Gaps
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0
                                   100.0%; Score 105; DB 2; Length 22; 100.0%; Pred. No. 9.2e-07; Live 0; Mismatches 0; Indels
sequence represents a peptide from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid encoding apoproteinA-I agonist peptides
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DUFOURCO J

(CORN/) (METZ/) (DUFO/)

SEKU/ BUTT/

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atherosclerosis, restenosis, HDL (high density lipoprotein) and apoA-I deficiency, hypertriglyceridemia and metabolic syndrome, also to treat endotoxemia (septic shock). Host cells containing (A) can also be used to study the role of apoA-I in lipid metabolism. (B) can be used diagnostically, e.g. to measure serum HDL (particularly its subpopulation involved in retrograde cholesterol transport) and for imaging the circulatory system or HDL accumulations at fatty streaks. The present sequence represents a peptide from the present invention
dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease.
                                                                                                                                                                                    Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-1999;
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(SEKU/)
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                                                                                                                                                                                                                                                           Matches
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                               The present invention describes a nucleic acid (A) encoding an apolipoprotein A-I (apoA-I) agonist (B) that is a peptide, or analog, which forms an amphipathic alpha-helix in presence of lipids. (A), optionally as a complex with lipids, and host cells that contain (A), are useful for gene therapy, or prevention, of diseases associated with captulation as pecifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) and apoA-I deficiency, hypertriglyceridemia and metabolic syndrome, also to treat cendotoxemia (septic shock). Host cells containing (A) can also be used to study the role of apoA-I in lipid metabolism. (B) can be used diagnostically, e.g. to measure serum HDL (particularly its subpopulation circulatory system or HDL accumulations at fatty streaks. The present circulatory system or HDL accumulations at fatty streaks. The present sequence represents a peptide from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gene therapy; apolipoprotein A-I; agonist; dyslipidemic disorder; ApoA-I; cardiovascular disease; atherosclerosis; restenosis; LCAT; hyperlipidemia; septic shock; lecithin:cholesterol acyltransferase.
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100.0%; Score 105; DB 2; Length 22;
Best Local Similarity 100.0%; Pred. No. 9.2e-07;
Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Metz G,
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                                                                                                                                                                                                                                                                                                                                                                                                                          1 PVLDLFRELLNELLEALKOKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PVLDLFRELLNELLEALKOKLK 22
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 Example; Page 157; 232pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention describes an apolipoprotein A-I (ApoA-I) agonist (A) comprising a 15-29 residue peptide or peptide analogue which forms an amphipathic alpha-helix in the presence of lipids or its salt. (A) is useful for treating a subject suffering from a disorder associated with dyslipidaemia (hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, high density lipoprotein (HDL) or Apia-I deficiency, hypertriglyceridaemia or metabolic syndrome) or septic shock. This is the amino acid sequence of a ApoA-I agonist peptide.
                                                                                                                                                                                                                                                                                               apolipoprotein A-I; ApoA-I; agonist; peptide analogue; amphipathic alpha-helix; dyslipidaemia; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; high density lipoprotein; HDi, Apia-I deficiency; hypertriglyceridaemia; metabolic syndrome; septic shock.
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100.0%; Score 105; DB 2; Length 22; 100.0%; Pred. No. 9.2e-07;
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Mismatches
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                                                               1 PVLDLFRELLNELLEALKOKLK 22
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                                22; Conservative
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BUTTNER K.
CORNUT I.
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                 Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes an apolipoprotein A-I (ApoA-I) agonist (A) comprising a 15-29 residue peptide or peptide analogue which forms an amphipathic alpha-helix in the presence of lipids or its salt. (A) is useful for treating a subject suffering from a disorder associated with dyslipidaemia (hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, high density lipoprotein (HDL) or Apia-I deficiency, hypertriglyceridaemia or metabolic syndrome) or septic shock. This is the amino acid sequence of a ApoA-I agonist peptide.
                                                                                                                                                                                                                                                                                                                                                  apolipoprotein A-I; ApoA-I; agonist; peptide analogue; amplipathic alpha-helix; dyslipidaemia; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; high density lipoprotein; HDL; Apia-I deficiency; hypertriglyceridaemia; metabolic syndrome; septic shock.
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                                            / Match 100.0%; Score 105; DB 8; Length 22; Local Similarity 100.0%; Pred. No. 9.2e-07; nes 22; Conservative 0; Mismatches 0; Indels
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                                                                                                              1 PVLDLFRELLNELLEALKOKLK 22
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DUFOURCO J.
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SEKUL R.
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Gaps

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1 PVLDLFRELLNELLEALKOKLK 22

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GenCore version 6.2.1
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OM protein - protein search, using sw model

Run on:

August 16, 2007, 22:42:55 ; Search time 39 Seconds (without alignments) 54.276 Million cell updates/sec

US-10-715-895A-4 105 1 PVLDLFRELLNELLEALKQKLK 22

Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Scoring table:

283416 segs, 96216763 residues Searched:

283416 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 80:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES		259	274 2 AI1256	274 2 AE1619 hypothetical	876 2 B96693 probable rece	225 1 E53402	577 2 T41727	86 2 B97107 hypothetical	195 2 JC8002 Clorf36 pi	276 2 G83527	325 2 F70440 conserved	303 2 H81395	344 2 C69453	478 2 JC5670	738 2 A53542	1082 1 RNEGB	179 2 G70325	242 2 D97229	291 2 C71877	317 2 G59103 hypothetical	379 2 AD1446 Protein gp4 (	414 2 E64387	438 2 AB1140 6-phospho-bet	548 2 G97047	566 2 E81404	327 2 E71130	199 2 E96926	335 2 C83927	379
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RESULT 2 A11256 hypothetical protein lmo1457 [imported] - Listeria monocytogenes (strain EGD-e)

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30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

## ALIGNMENTS

oprotein A-I precursor rat  130-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004  130-Jun-1988 #sequence_revision 30-Jun-1988 #text_changes S.K.  150-Jun-1988 #sequence_revision of the rat apolipoprotein A-I, C-III, and A-I, Chem. 261, 1266-1277, 1366  151-150-150-150-150-150-150-150-150-150-	 RESULT 1
C. Saccession: A24700, S00239; Moint and expression 30-Jun-1988 #text_change 09-Jul-2004 C. Saccession: A24700, S00239; Moint and Expression of the rat apolipoprotein A-1, C-III, and A-1 B. Hiddadd, I.A., Ordovas, J. M., Fitzpatrick, T.; Karathanasis, S.K. J. Hiddadd, I.A., Ordovas, J. M., Fitzpatrick, T.; Karathanasis, S.K. A; Title: Linkage, evolution, and expression of the rat apolipoprotein A-1, C-III, and A-1 A; Accession: A24700 A; Accession: A24700 A; A; Accession: A24700 A; A; Colouin, J.E.; Martial, J.A., Gialan, J.E. Bur. J. Biochem 140, 493-498, 1984 A; Foncin, J.E.; Martial, J.A., Gialan, J.E. Bur. J. Biochem 140, 493-498, 1984 A; Foncin, J.E.; Martial, J.A., Gialan, J.E. Bur. J. Biochem 140, 493-498, 1984 A; Foncin, J.E.; Martial, J.A., Gialan, J.E. Bur. J. Biochem 140, 493-498, 1984 A; Foncin, J.E.; Martial, J.A., Gialan, J.E. Bur. J. Biochem 140, 493-498, 1984 A; Foncin, J.E.; Martial, J.A., Gialan, J.E. A; Residues: 1.259 eRDN A; Residues: 1.250 eRDN A; Residu	 apolipoprotein A-I precursor - rat C;Species: Rattus norveqicus (Norway rat)
Windedd I.A., Ordovac J.M., Fitzpatrick, T.; Karathanasis, S.K.  S.Haddad, I.A., Ordovac J.M., Fitzpatrick, T.; Karathanasis, S.K.  J. BAO. Then. 261, 13268-1327, 1996 A; Recession. A24700 A; Recession. A24700 A; Recession. A24700 A; Residuce: 1.259 - GNDA A; Reference number: 500298; WUID:84207987; PMID:6426956 A; Recession: 500208 A; Residuce: 1.259 - GNDA A; Reference number: 500298; WUID:84207987; PMID:6426956 A; Residuce: 1.259 - GNDA A; Residuce: 1.250 - GNDA A; Reference number: Asidith, Dr.A. Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. A; Reference number: Asidith, Dr.A. Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. A; Reference number: Asidith, Dr.A. Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. A; Reference number: Asidith, Dr.A. Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. A; Residuce: 1.257, 971-971-978, 1932 A; Title: The primary translation product of rat intestinal apolipoprotein A; Residuce: UNIPARC:UPIO0017779 A; Residuce: 1.257, 971-971-978 A; Residuce: 1.247, Andy, R.; Alpers, A.; Alpers, A.	C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
A;Tille: Linkage, evolution, and expression of the rat apolipoprotein A-1, C-III, and A-1; A;Tille: Linkage, evolution, and expression of the rat apolipoprotein A-1, C-III, and A-1; A;Tille: Linkage, evolution, and expression of the rat apolipoprotein A-1. C-III, and A-1; A;Accession: A24700  A;Residues: 1-259 c:RAD  A;Residues: 1-250 c:RAD  A;	R.Haddad, 1.7A.; Ordovas, J.M.; Fitzpatrick, T.; Karathanasis, S.K. T. Biol Chem. 361 11368-11377 1986
Accession: A24700 A.Mccesslon: A24700 A.Mccesslon: A24700 A.Mccesslon: A24700 A.Mccesslon: A24700 A.Mccesslon: A24700 A.Mccesslon: A24704 A.Mccesslon: A259 cMAD A.Mccesslon: A259 cMAD A.Mccesslon: A259 cMAD A.Mccesslon: BATLall, J.A.; Glalen, J.E. BRIT. J. Biochem. 140, 439-498, 1984 A.Title: Cloning and structure analysis of the rat apolipoprotein A-I cDNA. A.Mcession: S00298; MUDD:84207987; PMID:6426956 A.Mccession: S00298; MUDD:84207987; PMID:6426956 A.Mccession: MAD A.Mccession: MAD A.Mccession: A259 cMAD A.Mccession: A251, 971-978, 1982 A.Mccession: A251, 971-978, 1982 A.Mccession: A2514 A.	 A; Title: Linkage, evolution, and expression of the rat apolipoprotein A-I, C-III, and A-1 A:Reference number: A92558; MUID:87008540; PMID:3020028
A; Residues: 1.259 cAND.  A; Cross-references: UNIPROT: P04639; UNIPARC: UPI000168276; EMBL: J02597; NID: g202935; PIDR R; Poncin, J.B.; Martial. J.A.; Gialen, J.E.  Bur. J. Biochem. 140, 493-498, 1984  A; Title: Cloning and structure analysis of the rat apolipoprotein A-I cDNA.  A; Reference number: S00298; MUD: 84207987; PMID: 6426956  A; Recence number: S00298; MUD: 84207987; PMID: 6426956  A; Residues: 1-259 cPON- A; Residues: 1-259 cPON- A; Cordon, J.I.; Smith, D.P.; Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. Biol. Chem. 257, 971-978, 1982 A; Ricerence number: A05314; MUD: 82098162; PMID: 6798036  A; Reference number: A05314; MUD: 82098162; PMID: 6798036 A; Rocesion: A05314 A; Wolecule type: protein A; Rocesion: A05314 A; MUD: 82098162; PMID: 6798036 A; Rocesion: A05314 A; Molecule type: protein A; Reference number: A05314 A; Molecule type: protein C; Comment: This protein is synthesized in the liver and small intestine. The propeptide C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Ruperfamily: apolipoprotein A-I C; Ruperfamily: Papolipoprotein A-I C; Rup	A;Accession: A24700 A;Molecule type: DNA
Bur. 0. Biochem. 140, 493-498, 1984  Ay Title: (Loning and structure analysis of the rat apolipoprotein A-I cDNA.  Ay Reference number: S00298, WUID:84207987; PMID:6426956  Ay Accession: S00298  Ay Accession: S00298  Ay Roccession: S00298  Ay Title: The primary translation product of rat intestinal apolipoprotein A-I mRNA is an Ay Reference number: AD314, WUID:82098162; PMID:6798036  Ay Roccession: AD314  Ay	A; Residues: 1-259 cray. A; Cross-references: UNIPROT: P04639; UNIPARC: UPI0000168276; EMBL: J02597; NID: 9202935; PIDP D: Donorin J F . Martial J A . Giolon J R
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A; Molecule type: mRNA A; Molecule type: mRNA A; Molecule type: mRNA A; Residues: 1-259 < PON> A; Molecule type: mRNA A; Residues: 1-259 < PON> A; Cosson references: UNIPARC: UP10000168276; GB: M00001; EMBL: X00558; NID: g202944; PIDN: AAAA; CGCAGON, J. I.; Smith, D. P.; Andy, R.; Alpers, D. H.; Schonfeld, G.; Strauss, A.W. J. Biol. Chem. 257, 971-979; 1982 A; Pride translation product of rat intestinal apolipoprotein A-I mRNA is an A; Reference number: A05314; MUD: 82098162; PMID: 6798036 A; Residues: 1-18, XX, 20-21, XX,	Alitte: Cloning and Structure analysis of the fat apolitybelouelin A-1 Colon. AlReference number: S00298; MUID:84207987; PMID:6426956
A; Residues: 1-259 <pon. 1-18,="" 1982="" 20-21,="" 23-28,="" 257,="" 30-38,="" 40-41,="" 43-45="" 971-978,="" <gor="" a-i="" a.w.="" a05314="" a05314;="" a;="" accession:="" alpers,="" an="" andy,="" apolipoprotein="" biol.="" chem.="" cross-references:="" d.h.;="" d.p.;="" embl:x00558;="" g.;="" gb:m00001;="" gordon,="" intestinal="" is="" j.="" j.i.;="" mena="" molecule="" muid:82098162;="" nid:g202944;="" number:="" of="" pidn:aaa,="" pmid:6798036="" primary="" product="" protein="" r.;="" r;="" rat="" reference="" residues:="" schonfeld,="" smith,="" strauss,="" the="" title:="" translation="" type:="" uniparc:upi000168276;="" x',=""> A; Cross-references: UNIPARC:UPI0000177798 C; Comment: This protein is synthesized in the liver and small intestine. The propeptide: C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Genetics: A; Interpretamily: apolipoprotein A-I C; Superfamily: apolipoprotein A-I C; Superfamily: apolipoprotein A-I C; Superfamily: apolipoprotein A-I #status experimental <ago. #status="" #status<="" 1-18="" 1-25-259="" <ago.="" a-i="" apolipoprotein="" domain:="" experimental="" f;="" product:="" propeptide="" th=""><th> A;Accession: S00298 A;Molecule type: mRNA</th></ago.></pon.>	 A;Accession: S00298 A;Molecule type: mRNA
R;Gordon, J.I.; Smith, D.P.; Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W. J. Biol. Chem. 257, 971-978, 1982  A;Title: The primary translation product of rat intestinal apolipoprotein A-I mRNA is an A;Reference number: A05314; MUID:82098162; PMID:6798036  A;Accession: A05314  A;Molecule type: protein A;Residues: 1-18, X', 20-21, X', 23-28, X', 30-38, X', 40-41, X', 43-45 <gor> A;Comment: This protein is synthesized in the liver and small intestine. The propeptide C;Comment: This protein is synthesized in the liver and small intestine in plasma. C;Genetics: A;Introns: 15/1; 66/2 C;Superfamily: apolipoprotein A-I C;Reywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol C;Reywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol F;19-18/Domain: signal sequence #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status ∏ A-I Manaches</pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></gor>	 A;Residues: 1-259 <pon> A:Residues: 1-259 <pon> A:Cross-references: INITARC:UPIO000168276; GB:M00001; EMBL;X00558; NID:q202944; PIDN:AAA4</pon></pon>
b Bloi. Chem. 257, 917-918, 1982  J. Bloi. Chem. 257, 917-918, 1982  A; Title: The primary translation packet of rat intestinal apolipoprotein A-I mRNA is an A; Reference number: A05314; MUID:82098162; PMID:6798036  A; Accession: A05314  A; Molecule type: protein  A; Residues: 1-18, 'X', 20-21, 'X', 23-28, 'X', 30-38, 'X', 40-41, 'X', 43-45 <gor> A; Cross-references: UNIPARC:UPT0000177798  C; Comment: This protein is synthesized in the liver and small intestine. The propeptide: C; Comment: This protein is a major component of the high density lipoproteins in plasma. C; Genetics: A; Introns: 15/1; 66/2  C; Superfamily: apolipoprotein A-I  C; Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipoprotein: signal sequence #status experimental <pro> F; 1-18/Domain: signal sequence #status experimental <pro> F; 1-14/Domain: propeptide #status experimental <pro> F; 1-14/Domain: propeptide #status experimental <pro> F; 1-14/Domain: propeptide #status experimental <pro> F; 25-259/Product: apolipoprotein A-I #status &amp;PRO&gt; F; 25-259/P</pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></gor>	 R; Gordon, J.I.; Smith, D.P.; Andy, R.; Alpers, D.H.; Schonfeld, G.; Strauss, A.W.
A;Reference number: A05314; MUID: B2098162; PMID: 6798036 A;Accession: A05314 A;Accession: A05314 A;Accession: A05314 A;Accession: A05314 A;Accession: A; X, 20-21, X', 23-28, X', 30-38, X', 40-41, X', 43-45 <gor> A;Cross-references: UnlPARC:UPID:00177798 A;Cross-references: UnlPARC:UPID:00177798 C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins A: A;Introns: 15.1 66/2 C;Superfamily: apolipoprotein A-I C;Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipoprofes in: propeptide #status experimental <pro> F;1-18/Domain: signal sequence #status experimental <pro> F;1-18/Domain: propeptide #status experimental <pro> F;1-18/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <ando #status="" <ando="" a-i="" apo<="" apolipoprotein="" experimental="" f;25-259="" product:="" th=""><th>J. BIOI. CREM. 25/, 9/1-9/8, 1982 A;Title: The primary translation product of rat intestinal apolipoprotein A-I mRNA is an</th></ando></pro></pro></pro></pro></gor>	J. BIOI. CREM. 25/, 9/1-9/8, 1982 A;Title: The primary translation product of rat intestinal apolipoprotein A-I mRNA is an
A,Accession: A02114 A,Accession: A02114 A,Accession: A02114 A,Accession: A021, X', 23-28, X', 30-38, X', 40-41, X', 43-45 <gor> A,Cross-references: Uniparacipation (1779) A,Cross-references: Uniparacipation (1779) C,Comment: This protein is synthesized in the liver and small intestine. The propeptide (2,Comment: This protein is a major component of the high density lipoproteins in plasma. C,Genetics: C,Comment: This protein is a major component of the high density lipoproteins in plasma. C,Genetics: A,Introns: 15/1, 66/2 C,Superfamily: apolipoprotein A-I C,Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipop C,Superfamin: signal sequence #status experimental <pro> F;19-24/Domain: signal sequence #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <and 0;="" 122="" 12;="" 143<="" 22="" 259;="" 2;="" 4;="" 54.5*;="" 6;="" 8.3;="" best="" c,="" conservative="" db="" dd="" gaps="" guery="" i="" indels="" length="" local="" match="" matches="" mismatches="" no.="" phldepoekwhendryprayrqkle="" pred.="" pvldlfrblinellerianckik="" s3;="" score="" similarity="" so.s*;="" th=""><th>A; Reference number: A05314; MUID: 82098162; PMID: 6798036</th></and></pro></pro></pro></pro></gor>	A; Reference number: A05314; MUID: 82098162; PMID: 6798036
A;Residues: 1-18,7x',20-21,7x',23-28,7x',30-38,7x',40-41,7x',43-45 <gor> A;Cross-references: UnlPRC:UPID0017798 C;Comment: This protein is synthesized in the liver and small intestine. The propeptide c;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: Signal is metabolism; HDL; intestine; lipid binding; lipid transport; lipop C;Superfamily: apolipoprotein A-I #status experimental <pro> F;1-18/Domain: signal sequence #status experimental <pro> F;1-18/Domain: propeptide #status experimental <pro> F;1-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <mat> Cuery Match</mat></pro></pro></pro></pro></gor>	 A;Accession: A05314 A:Wolecule type: protein
A;Cross-releated so unity that C: UTUOUL/1/36  C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Gonetics:  C;Gonetics:  C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Comment: 15.1  C;Gonetics:  C;Comment: 11, 66/2  C;Superfamily: apolipoprotein A-I  C;Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipoprofe; lipoprotein: signal sequence #status experimental <pro>  F;1-18/Domain: propeptide #status experimental <pro> F;1-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <mat>  Cuery Match  So.5*; Score S3; DB 2; Length 259; Best Local Similarity 54.5*; Pred. No. 8.3;  Matches 12; Conservative 4; Mismatches 6; Indels 0; Gaps 0;  Oy I PVLDEPRELIMELLERIANCILES (1)                                      </mat></pro></pro></pro>	 A;Residues: 1-18, X,,20-21, X,,23-28, X,,30-38, X,,40-41, X,,43-45 <gor></gor>
C;Comment: This protein is a major component of the high density lipoproteins in plasma. C;Gonetics A;Introns: 15/1; 66/2 C;Superfamily: apolipoprotein A-I C;Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol C;Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol C;Keywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol C;Keywords: chomain: signal sequence #status experimental <pro> F;1-18/Domain: signal sequence #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <ato 0;="" 122="" 12;="" 143<="" 22="" 259;="" 2;="" 4;="" 54.5%;="" 6;="" 8.3;="" best="" chery="" conservative="" db="" gaps="" i="" indels="" length="" local="" match="" matches="" mismatches="" no.="" phldepoekwheneryroxie="" plidepoekwhenerneryroxik="" pred.="" pvldeprelimelleriancian="" qy="" s3;="" score="" similarity="" so.5%;="" th=""><th>A;Cross-references: UNIPARC:UP10000177798 C;Comment: This protein is synthesized in the liver and small intestine. The propeptide i</th></ato></pro></pro>	A;Cross-references: UNIPARC:UP10000177798 C;Comment: This protein is synthesized in the liver and small intestine. The propeptide i
C;Genetics 15/1; 66/2 C;Superfamily: apolipoprotein A-I C;Superfamily: apolipoprotein A-I C;Superfamily: apolipoprotein A-I C;Superfamily: apolipoprotein A-I C;Seywords: cholesterol metabolism; HDL; intestine; lipid binding; lipid transport; lipol C;Seywords: chomain: signal sequence #status experimental <pro> F;1-18/Domain: signal sequence #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <ai #status="" 0;="" 12;="" 4;="" 6;="" <ai="" a-i="" apolipoprotein="" chery="" conservative="" experimental="" f;25-259="" gaps="" i="" indels="" match="" matches="" mismatches="" oy="" product:="" pvldeprelimententententententententententententente<="" th=""><th>C; Comment: This protein is a major component of the high density lipoproteins in plasma.</th></ai></pro></pro></pro></pro>	C; Comment: This protein is a major component of the high density lipoproteins in plasma.
C;Superfamily: apolipoprotein A-I C;Savperfamily: apolipoprotein A-I C;Kavpordas: cholesterol metabolism, HDL; intestine; lipid binding; lipid transport; lipol C;Kavpordas: cholesterol metabolism, HDL; intestine; lipid binding; lipid transport; lipol C;Kavpordas: signal sequence #status experimental <pro> F;19-24/Domain: signal sequence #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <pro> R;25-259/Product: apolipoprotein A-I #status experimental <pro #status="" <pro="" a-i="" apolipoprotein="" experimental="" product:="" production="" r;25-259="" r;25<="" th=""><th>C;Genetics: A;Introns: 15/1; 66/2</th></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro></pro>	C;Genetics: A;Introns: 15/1; 66/2
F;1-18/Domain: signal sequence #status experimental <pgs #status="" <pro="" domain:="" experimental="" f;19-24="" propeptide=""> F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <mat> Query Match</mat></pro></pgs>	 C, Superfamily: apolipoprotein A-I C. Roxwords: cholesterol metaholism: HDL: intestine: libid binding: libid transport: libor
F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <pat> Query Match  Goury Match  Best Local Similarity 54.5%; Pred. No. 8.3;  Matches 12; Conservative 4; Mismatches 6; Indels 0; Gaps 0;  Qy</pat></pro>	Fil-18/Domain: signal sequence experimental Sign
o; Gaps	 F;19-24/Domain: propeptide #status experimental <pro> F;25-259/Product: apolipoprotein A-I #status experimental <mat></mat></pro>
3est Local Similarity 54.5%; Pred. No. 8.3;  Aatches 12; Conservative 4; Mismatches 6; Indels 0; Gaps  1 PVLDLFRELLNELLEALKQKLK 22	50.5%; Score 53; DB 2;
	54.5%; pred. No. 8.3; ative 4; Mismatches 6; Indels 0; Gaps

us-10-715-895a-4.rpr

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Gaps

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A; Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C.: C.A.; Li, J.H.; Li, Y.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A; Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, I ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A; Fitle: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A; Reference number: A86141; MUID:21016719; PMID:11130712
A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                serine O-acetyltransferase (EC 2.3.1.30) cysE [similarity] - Bacillus stearothermophilus C; Species: Bacillus stearothermophilus C; Species: Bacillus stearothermophilus C; Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 05-May-2000 C; Accession: E53402 R; Gagnon, Y: Breton, R.; Putzer, H.; Pelchat, M.; Grunberg-Manago, M.; Lapointe, J. J. Biol. Chem. 269, 7473-7482, 1994 A; Title: Clustering and co-transcription of the Bacillus subtilis genes encoding the amir
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                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q9C622; UNIPARC:UP1000004840F; GB:AE005173; NID:g11128393; P1
C;Ganetics:
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C,Function:
A,Fathway: cysteine biosynthesis
A,Note: rate-limiting step
C,Superfamily: Bacillus serine acetyltransferase; serine acetyltransferase homology
C,Keywords: acyltransferase; coenzyme A
F;9-169/Domain: serine acetyltransferase homology <SAT>
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C;Species: Schizosaccharomyces pombe
C;Species: Schizosaccharomyces pombe
C;Accession: T41727
R;Lyne, M.; Wood, V.; Rajandream, M.A.; Barrell, B.G.; Murphy, L.; Harris, D submitted to the EMBL Data Library, June 1998
A;Reference number: Z22013
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9
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A;Accession: B53402
A;Status: preliminary; not compared with conceptual translation
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A;Residues: 1-225 <GAG>
A;Cross-references: UNIPARC:UP10000172390
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |:|| || :|| || || 388 PLLDRFRGVLNNLREMCKRKV 408
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les 13; Conserv
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A; Residues: 1-876 <STO>
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C;Species: Listeria innocua
C;Species: Listeria innocua
C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
C;Accession: AE1619
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker, Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi, H. Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, A.; Ma A;Atutle: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Status: preliminary
A;Nolecule type: DNA
A;Residues: 1-274 <GLA>
C;Species: Listeria monocytogenes
C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
C;Jaccession. A11256
C;Accession. A1126
C;Accession. B. Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker
C;Accession. A1126
B;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker
C; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi, H.
D; Jones, L.M.; Karst, J.
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, A.; Ma
A;Authors: Kreft, J.; Sinoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; Wehland,
A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
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A,Residues: 1-274 <GLA>
A;Cross-references: UNIPROT:092BQ3; UNIPARC:UPI00000553A4; GB:NC_003210; PIDN:CAC99535.1
A;Experimental source: strain EGD-e
C;Gene: Lactics:
A;Gene: lmo1457
C;Superfamily: ydiA protein
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B96693
By Cobable receptor serine/threonine kinase PRSK T4024.2 [imported] - Arabidopsis thaliana C; Species: Arabidopsis thaliana (mouse-ear cress)
C; Species: Arabidopsis thaliana (mouse-ear cress)
C; Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C; Accession: B96693
R; Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; Anture 408, 816-820, 2000
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8.8;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50.5%; Score 53; 55.0%; Pred. No.
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C,Genetics:
A,Gene: lin1494
C,Superfamily: ydiA protein
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Best Local Similarity
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Best Local Similarity
Matches 11; Conserv
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C;Species: Pseudomonas aeruginosa
C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
C;Accession: G83527
R;Stover, C:K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Briadman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, J. Lory, S.; Olson, M.V.
Nature 406, 595-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathog A,Recession: G83527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      conserved hypothetical protein ag_1630 - Aquifex aeolicus
C;Species: Aquifex aeolicus
C;Dates: 08-May-1998 #sequence_revision 08-May-1998 #text_change 09-Jul-2004
C;Accesion: F70440
R;Deckert, G; Warren, P.V.; Gaasterland, T.; Young, W.G.; Lenox, A.L.; Graham, D.E.; Ove
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C;Species: Campylobacter jejuni
C;Species: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 09-Jul-2004
C;Accession: H81395
C;W.; Quail, W.; Wren, B.W.; Mungall, K.; Ketley, J.M.; Churcher, C.; Basham, D.; Chilling C:W.; Quail, M.; Rajandream, M.A.; Rutherford, K.M.; VanVliet, A.; Whitehead, S.; Barrel)
Nature 403, 665-668, 2000
A;Title: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hype
A;Reference number: A81250; MuID:20150912; PMID:10688204
                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: DNA
A;Residues: 1.276 <53TO>
A;Cross-references: UNIPROT:091523; UNIPARC:UP100000C51C7; GB:AE004528; GB:AE004091; NID:
A;Experimental source: strain PAO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Gene: PA0935
C;Superfamily: nucleoside triphosphate pyrophosphohydrolase MazG; beta-lactamase regulato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:067552; UNIPARC:UPI0000056687; GB:AE000748; NID:g2983960; PII A;Experimental source: strain VP5
                      conserved hypothetical protein PA0935 [imported] - Pseudomonas aeruginosa (strain PAO1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 392, 353-358, 1998
A;Title: The complete genome of the hyperthermophilic bacterium Aquifex aeolicus. A;Title: The complete genome of the hyperthermophilic bacterium Aquifex aeolicus. A;Reference number: A70300; MUID:9819666; PMID:9537320
A;Accession: P70440
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
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C;Superfamily: Mycoplasma conserved hypothetical protein MG190
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176 PVVDKVREGLDEVLEAMAE 194
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303 YERALKKULEAIKBELK 319
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nes 9; Conserv
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A;Cross-references: GB:AY191519
C;Comment: This protein is a evolutionary conserved phosphorylated protein that has a mi
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C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004
C;Accession: B97107
R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee, J. Daly, M.J.; Bennett, G.N.; Koonin, B.V.; Smith, D.R.
J. Bacteriol. 183, 4833-4838, 2001
A;Reference and Comparative Analysis of the Solvent-Producing Bacterium Clc A;Reference number: A96900; MUID:21359325; PMID:21359325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A,Cross-references: UNIPROT:0971G2; UNIPARC:UP100000CA2A9; GB:AE001437; PIDN:AAK79645.1; A;Experimental source: Clostridium acetobutylicum ATCC824
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C;Accession: JC8802

R;Lavorgna, G.; Lestingi, M.; Ziviello, C.; Testa, F.; Simonelli, F.; Manitto, M.P.; Br

Biochem. Biophys. Res. Commun. 308, 414-421, 2003

A;Title: Identification and characterization of Clorf36, a transcript highly expressed

A;Reference number: JC8802; PMID:12914764
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C;Species: Homo sapiens (man)
C;Date: 10-Nov-2003 #sequence_revision 10-Nov-2003 #text_change 24-Nov-2003
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Pred. No. 21;
4; Mismatches 5; Indels
                                                                                                                                                 Query Match

48.6%; Score 51; DB 2; Length 577;

Best Local Similarity 57.1%; Pred. No. 35;

Matches 12; Conservative 3; Mismatches 6; Indels
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C;Keywords: Clorf36; coiled-coil; mitochondrial function
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Pred. No. 9.2;
5; Mismatches
A;Experimental source: strain 972h-; cosmid c338 C;Genetics: A;Gene: pof3; SPDB:SPCC338.16 A;Map position: 3
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Best Local Similarity 52.6%;
Matches 10; Conservative
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Best Local Similarity 52.9%;
Matches 9; Conservative
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A;Molecule type: DNA
A;Residues: 1-86 <KUR>
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RESULT 8

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RESULT 9

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Search completed: August 16, 2007, 22:49:16
Job time : 42 secs
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C;Keywords: Golgi apparatus
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-738 <POD>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Accession: S09210
A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                          C; Genetics:
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 05-Dec.1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession. G59453.
R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson
J; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.
Nature 390, 364-370, 1997
A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S.
Smith, H.O.; Woese, C.R.; Venter, J.C.
A;Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archaec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA

A; Moelecule type: mRNA

A; Residues: 1-478 c.TAb.

A; Cross-references: UNIPROT: 035114; UNIPARC: UPI00000231D6; GB: AB008553; NID: 92618485; PI

C; Comment: This protein is involved in sequestration of particular cytoplasmic proteins

C; Superfamily: lysosomal integral membrane protein II

F; 1-26/Domain: signal sequence #status predicted cSIG>

F; 433-458/Domain: transmembrane #status predicted cYPM>

F; 474-475/Region: endosomal/lysosomal sorting signal
                    A;Residues: 1.303 <PAR>
A;Residues: 1.303 <PAR>
A;Cross-references: UNIPROT:Q9PI08; UNIPARC:UPI000012C5CC; GB:AL139075; GB:AL111168; NID
A;Experimental source: serotype O2, strain NCTC 11168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:028645; UNIPARC:UPI000056C62; GB:AE000990; GB:AE000782; NID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Tabuchi, N.; Akasaki, K.; Sasaki, T.; Kanda, N.; Tsuji, H.
D. Biochman. 122, 756-763, 1997
A;Title: Identification and characterization of a major lysosomal membrane glycoprotein,
A;Reference number: JC5670; MUID:98060500; PMID:9399579
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Residues: 1-344 <KLE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lysosomal membrane 85K glycoprotein precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 20-Nov-1997 #sequence_revision 20-Nov-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                              Score 47; DB 2;
Pred. No. 62;
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Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 LDLFRELLNELLEALKOKL 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               267 YŘKĽVRKĽFEGĽKÓNĽK 283
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                                                                                                                                                                                           Query Match
Best Local Similarity 47.4%;
Matches 9; Conservative
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Best Local Similarity
Free 9; Conserve
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Best Local Similarity
Matches 10; Conserv
                                                                                                                    A;Gene: Cj0501; Cj0503c
C;Keywords: lyase
A; Molecule type: DNA
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                                                                                             C, Genetics:
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Drefeldin A-sensitive Golgi protein LDLC - human C.Species: Homo sapiens (man)
C.Species: Landom sapiens (man)
C.Date: 25-Aug-1995 #sequence_revision 25-Aug-1995 #text_change 09-Jul-2004
C.Date: 25-Aug-1995 #sequence_revision 25-Aug-1995 #text_change 09-Jul-2004
C.Accession: A53542
R.Podos, S.D.; Reddy, P.; Ashkenas, J.; Krieger, M.
J. Cell Biol. 127, 679-691, 1994
A.Fitle: LDLC encodes a brefeldin A-sensitive, peripheral Golgi protein required for norm A.Reference number: A53542; MUID:95050941; PMID:7962052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA-directed RNA polymerase (EC 2.7.7.6) beta chain - Euglena gracilis chloroplast C; Species: chloroplast Euglena gracilis C; Species: chloroplast Euglena gracilis C; Species: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004 C; Accession: S03210 R; Radebaugh, C.A.; Hallick, R.B. Nucleic Acids Res: 18, 1865-1878, 1990 A; Fitel: The Euglena gracilis chloroplast rpoB gene. Novel gene organization and transcri A; Reference number: S09210; MUID:90245579; PMID:2110656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-1082 <YEP>
A;Cross-references: UNIPROT:P23579; UNIPARC:UPI00001346D7; EMBL:X17191; NID:g11501; PIDN: A;Note: the authors translated the codon CGT for residue 132 as Gly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:Q14746; UNIPARC:UP10000127E38; GB:Z34975; NID:g575653; PIDN:C
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A;Introns: 137,3; 89/2; 117/1; 154/3; 227/3; 241/1; 288/2; 732/3
C;Superfamily: DNA-directed RNA polymerase beta chain
C;Keywords: chloroplast; nucleotidyltransferase; transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 738;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Pred. No. 1.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 LDLFRELLNELLEALKOKLK 22
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# Search Report

#### STIC Database Tracking Number: 234005

To: MARK SHIBUYA

Location: REM-2A29 / Mailbox 2C18

Art Unit: 1639

Thursday, August 16, 2007

Case Serial Number: 10/715895

From: HUYEN-TRAN TON-NU

Location: EIC 1600 REM-1D58 / REM-1B61 Phone: (571)272-9218

huyen-tran.ton-nu@uspto.gov

#### **Search Notes**

Dear Examiner SHIBUYA:

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services.

Huyen-Tran Ton-nu Technical Information Specialist STIC Biotech/Chem Library (571) 272-9218

